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#### (54) Primers for synthesising full-length cDNA and their use

(57) Primers for synthesizing full-length cDNAs and their use are provided.

5602 cDNA encoding a human protein has been isolated and nucleotide sequences of 5'-, and 3' -ends of the cDNA have been determined. Furthermore, prim-

ers for synthesizing the full-length cDNA have been provided to clarify the function of the protein encoded by the cDNA. The full-length cDNA of the present invention containing the translation start site provides information useful for analyzing the functions of the protein.

#### Description

#### FIELD OF THE INVENTION

The present invention relates to a polynucleotide encoding a novel protein, a protein encoded by the polynucleotide, and new uses of these.

#### BACKGROUND OF THE INVENTION

10 [0002] Currently, the sequencing projects, the determination and analysis of the genomic DNA of various living organisms have been in progress all over the world. The whole genomic sequences of more than 10 species of prokaryotes, a lower eukaryote, yeast, and a multicellular eukaryote, C. elegans are already determined. As to human genome, which is supposed to be composed of three thousand million base pairs, the world wide cooperative projects have been under way to analyze it, and the whole structure is predicted to be determined by the years 2002-2003. The aim of the determination of genomic sequence is to reveal the functions of all genes and their regulation and to understand living organisms as a network of interactions between genes, proteins, cells or individuals through deducing the information in a genome, which is a blueprint of the highly complicated living organisms. To understand living organisms by utilizing the genomic information from various species is not only important as an academie subject, but also socially significant from the viewpoint of industrial application.

[0003] However, determination of genomic sequences itself cannot identify the functions of all genes. For example, as for yeast, only the function of approximately half of the 6000 genes, which is predicted based on the genomic sequence, was able to be deduced. As for human, the number of the genes is predicted to be approximately one hundred thousand. Therefore, it is desirable to establish "a high throughput analysis system of the gene functions" which allows us to identify rapidly and efficiently the functions of vast amounts of the genes obtained by the genomic sequencing.

[0004] Many genes in the eukaryotic genome are split by introns into multiple exons. Thus, it is difficult to predict correctly the structure of encoded protein solely based on genomic information. In contrast, cDNA, which is produced from mRNA that lacks introns, encodes a protein as a single continuous amino acid sequence and allows us to identify the primary structure of the protein easily. In human cDNA research, to date, more than one million ESTs (Expression Sequence Tags) are publicly available, and the ESTs presumably cover not less than 80% of all human genes.

[0005] The information of ESTs is utilized for analyzing the structure of human genome, or for predicting the exonregions of genomic sequences or their expression profile. However, many human ESTs have been derived from proximal regions to the 3'-end of cDNA, and information around the 5'-end of mRNA is extremely little. Among these human cDNAs, the number of the corresponding mRNAs whose encoding protein sequences are deduced is approximately 7000, and further, the number of full-length therein is only 5500. Thus, even including cDNA registered as EST, the percentage of human cDNA obtained so far is estimated to be 10-15% of all the genes.

[0006] It is possible to identify the transcription start site of mRNA on the genomic sequence based on the 5'-end sequence of a full-length cDNA, and to analyze factors involved in the stability of mRNA that is contained in the cDNA, or in its regulation of expression at the translation stage. Also, since a full-length cDNA contains ATG, the translation start site, in the 5'-region, it can be translated into a protein in a correct frame. Therefore, it is possible to produce a large amount of the protein encoded by the cDNA or to analyze biological activity of the expressed protein by utilizing an appropriate expression system. Thus, analysis of a full-length cDNA provides valuable information which complements the information from genome sequencing. Also, full-length cDNA clones that can be expressed are extremely valuable in empirical analysis of gene function and in industrial application.

[0007] Therefore, if a novel human full-length cDNA is isolated, it can be used for developing medicines for diseases in which the gene is involved. The protein encoded by the gene can be used as a drug by itself. Thus, it has great significance to obtain a full-length cDNA encoding a novel human protein.

[0008] In particular, human secretory proteins or membrane proteins would be useful by itself as a medicine like tissue plasminogen activator (TPA), or as a target of medicines like membrane receptors. In addition, genes for signal transduction-associated proteins (protein kinases, etc.), glycoprotein-associated proteins, transcription-associated proteins, etc. are genes whose relationships to human diseases have been elucidated. Moreover, genes for disease-associated proteins form a gene group rich in genes whose relationships to human diseases have been elucidated.

[0009] Therefore, it has great significance to isolate novel full-length cDNA clones of human, only few of which has been isolated. Especially, isolation of a novel cDNA clone encoding a secretory protein or membrane protein is desired since the protein itself would be useful as a medicine, and also the clones potentially include a gene associated with diseases. In addition, genes encoding proteins that are associated with signal transduction, glycoprotein, transcription, or diseases are expected to be useful as target molecules for therapy, or as medicines themselves. These genes form a gene group predicted to be strongly associated with diseases. Thus, identification of the full-length cDNA clones

encoding those proteins has great significance.

#### SUMMARY OF THE INVENTION

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- 5 [0010] An objective of the present invention is to provide a polynucleotide encoding a novel protein, a protein encoded by said polynucleotide, and novel usages of these.
  - [0011] The inventors have developed a method for efficiently cloning a human full-length cDNA that is predicted by the ATGpr etc. to be a full-length cDNA clone, from a full-length-enriched cDNA library that is synthesized by the oligocapping method. Then, the inventors determined the nucleotide sequence of the obtained cDNA clones from both 5'- and 3'- ends.
  - [0012] Furthermore, the inventors analyzed the obtained clones by the BLAST search of the databases, SwissProt (http://www.ebi.ac.uk/ebi\_docsSwissProt\_db/swisshome.html), GenBank (http://www.ncbi.nlm.nih.gov/web/GenBank), and UniGene (Human) (http://www.ncbi.nlm.nih.gov/UniGene).
  - [0013] The full-length cDNA clones of the present invention have high fullness ratio since these were obtained by the combination of (1) construction of a full-length-enriched cDNA library that is synthesized by the oligo-capping method, and (2) a system in which the full-length ratio is evaluated from the nucleotide sequence of the 5'-end (selection based on the ATGpr, previously removed complete sequences to ESTs). However, the primer of the present invention enables to obtain full-length cDNA easily without any specialized methods as in the described method.
- Homology analysis in which the analysis is carried out against a not-full-length cDNA fragment to postulate the function of a protein encoded by said fragment, is being commonly performed.
  - However, since such analysis is based on the information of the fragment, it is not clear as to whether this fragment corresponds to a part that is functionally important in the protein. In other words, the reliability of the homology analysis based on the information of a fragment is doubtful, as information related to the structure of the whole protein is not available. However, the homology analysis of the present invention is conducted based on the information of a full-length cDNA comprising the whole coding region of the cDNA, and therefore, the homology of various portions of the protein can be analyzed. Hence, the reliability of the homology analysis has been dramatically improved in the present invention.
  - [0014] The inventors completed the invention by finding that it is possible to synthesize a novel full-length cDNA by using the combination of a primer that is designed based on the nucleotide sequence of the 5'-ends of the selected full-length cDNA clones and any of an oligo-dT primer or a 3'-primer that is designed based on the nucleotide sequence of the 3'-ends of the selected clones.
  - [0015] Thus, the present invention relates to primers described below, a method for synthesizing a polynucleotide using the primers, and polynucleotides obtained by the method.
  - [0016] First, the present invention relates to
    - (1) use of an oligonucleotide as a primer for synthesizing the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, or the complementary strand thereof, wherein said oligonucleotide is complementary to said polynucleotide or the complementary strand thereof and comprises at least 15 nucleotides;
    - (2) a primer set for synthesizing polynucleotides, the primer set comprising an oligo-dT primer and an oligonucleotide complementary to the complementary strand of the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, wherein said oligonucleotide comprises at least 15 nucleotides; and
    - (3) a primer set for synthesizing polynucleotides, the primer set comprising a combination of an oligonucleotide comprising a nucleotide sequence complementary to the complementary strand of the polynucleotide comprising a 5'-end nucleotide sequence and an oligonucleotide comprising a nucleotide sequence complementary to the polynucleotide comprising a 3'-end nucleotide sequence, wherein said oligonucleotides comprise at least 15 nucleotides and wherein said combination of 5'-end nucleotide sequence / 3'-end nucleotide sequence is selected from the combinations of 5'-end nucleotide sequence / 3'-end nucleotide sequence set forth in the SEQ ID NOs in Tables 1 and 2.
  - [0017] Tables 1 and 2 shows names of clones obtained in the examples described later, comprising the polynucleotide of the present invention (Table 1; 5547 clones, Table 2; 54 clones), names of nucleotide sequences at the 5'-end and 3'-end of the full-length cDNA, and their corresponding SEQ ID NOs. A blank indicates that the 3'-end sequence corresponding to the 5'-end sequence has not been determined for the same clone.
  - [0018] The SEQ ID NO of a 5'-end sequence is shown on the right side of the name of the 5'-end sequence, and the SEQ ID NO of a 3'-end sequence is shown on the right side of the name of the 3'-end sequence.

Table 1

	name of	SEQ ID	name of	SEQ ID
clone	5'-end	of 5'-end	3'-end	of 3'-end
***	sequence	sequence	sequence	sequence
HEMDA100000	E 11E1004100000	5 1	R-HEMBA10000	05 5548
HEMBA1000005	F-HEMBA100000	-	K-UEMDA 10000	05 5546
HEMBA1000012	F-HEMBA100001			
HEMBA1000020	F-HEMBA100002			
HEMBA1000030	F-HEMBA100003	0 4	R-HEMBA10000	30 5549
HEMBA1000042	F-HEMBA100004	2 5	R-HEMBA10000	42 5550
HEMBA1000046	F-HEMBA100004	6 6	R-HEMBA10000	46 5551
HEMBA1000050	F-HEMBA100005	0 7	R-HEMBA10000	50 5552
HEMBA1000076	F-HEMBA100007	6 8	R-HEMBA10000	76 5553
HEMBA1000111	F-HEMBA100011	1 9	R-HEMBA10001	11 5554
HEMBA1000129	F-HEMBA100012	9 10	R-HEMBA10001	29 5555
HEMBA1000141	F-HEMBA100014	1 11	R-HEMBA10001	41 5556
HEMBA1000150	F-HEMBA100015	0 12	R-HEMBA10001	50 5557
HEMBA1000156	F-HEMBA100015	6 13	R-nnnnnnnnn	nn 5558
HEMBA1000158	F-HEMBA100015	8 14	R-HEMBA10001	58 5559
HEMBA1000168	F-HEMBA100016	8 15	R-nnnnnnnnn	nn 5560

	HEMBA1000180	F-HEMBA1000180	16	R-HEMBA1000180	5561
	HEMBA1000185	F-HEMBA1000185	17	R-HEMBA1000185	5562
	HEMBA1000193	F-HEMBA1000193	18	R-HEMBA1000193	5563
5	HEMBA1000201	F-HEMBA1000201	19	R-HEMBA1000201	5564
	HEMBA1000213	F-HEMBA1000213	20	R-HEMBA1000213	5565
	HEMBA1000216	F-HEMBA1000216	21	R-HEMBA1000216	5566
	HEMBA1000227	F-HEMBA1000227	22	R-nnnnnnnnnnnn	5567
10	HEMBA1000231	F-HEMBA1000231	23	R-HEMBA1000231	5568
	HEMBA1000243	F-HEMBA1000243	24	R-HEMBA1000243	5569
	HEMBA1000244	F-HEMBA1000244	25	R-HEMBA1000244	5570
	HEMBA1000251	F-HEMBA1000251	26	R-HEMBA1000251	5571
15	HEMBA1000264	F-HEMBA1000264	27	R-HEMBA1000264	5572
	HEMBA1000280	F-HEMBA1000280	28	R-nnnnnnnnnnn	5573
	HEMBA1000282	F-HEMBA1000282,	29	R-HEMBA1000282	5574
	HEMBA1000288	F-HEMBA1000288	30	R-HEMBA1000288	5575
20	HEMBA1000290	F-HEMBA1000290	31	R-HEMBA1000290	5576
	HEMBA1000302	F-HEMBA1000302	32	R-HEMBA1000302	5577
	HEMBA1000303	F-HEMBA1000303	33	R-nnnnnnnnnnn	5578
	HEMBA1000304	F-HEMBA1000304	34	R-nnnnnnnnnn	5579
25	HEMBA1000307	F-HEMBA1000307	35	R-HEMBA1000307	5580
25	HEMBA1000327	F-HEMBA1000327	36		
	HEMBA1000333	F-HEMBA1000333	37	R-nnnnnnnnnnn	5581
	HEMBA1000338	F-HEMBA1000338	38	R-HEMBA1000338	5582
	HEMBA1000351	F-HEMBA1000351	39	R-HEMBA1000351	5583
30	HEMBA1000355	F-HEMBA1000355	40	R-HEMBA1000355	5584
	HEMBA1000356	F-HEMBA1000356	41		
	HEMBA1000357	F-HEMBA1000357	42	R-HEMBA1000357	5585
	HEMBA1000366	F-HEMBA1000366	43	R-HEMBA1000366	5586
35	HEMBA1000369	F-HEMBA1000369	44	R-HEMBA1000369	5587
	HEMBA1000376	F-HEMBA1000376	45	R-HEMBA1000376	5588
	HEMBA1000387	F-HEMBA1000387	46	R-HEMBA1000387	5589
	HEMBA1000390	F-HEMBA1000390	47	R-HEMBA1000390	5590
40	HEMBA1000392	F-HEMBA1000392	48	R-HEMBA1000392	5591
	HEMBA1000396	F-HEMBA1000396	49	R-HEMBA1000396	5592
	HEMBA1000411	F-HEMBA1000411	50	R-HEMBA1000411	5593
	HEMBA1000418	F-HEMBA1000418	51	R-HEMBA1000418	5594
45	HEMBA1000422	F-HEMBA1000422	52	R-HEMBA1000422	5595
	HEMBA1000428	F-HEMBA1000428	53	R-HEMBA1000428	5596
	HEMBA1000434	F-HEMBA1000434	54	R-HEMBA1000434	5597
	HEMBA1000442	F-HEMBA1000442	55	R-HEMBA1000442	5598
50	HEMBA1000456	F-HEMBA1000456	56	R-HEMBA1000456	5599
	HEMBA1000459	F-HEMBA1000459	57	R-HEMBA1000459	5600
	HEMBA1000460	F-HEMBA1000460	58	R-HEMBA1000460	5601
	HEMBA1000464	F-HEMBA1000464	59	R-HEMBA1000464	5602
55	HEMBA1000469	F-HEMBA1000469	60	R-HEMBA1000469	5603
JJ	HEMBA1000488	F-HEMBA1000488	61	R-HEMBA1000488	5604

	HEMBA1000490	F-HEMBA1000490	62	R-HEMBA1000490	5605
	HEMBA1000491	F-HEMBA1000491	63	R-HEMBA1000491	5606
	HEMBA1000501	F-HEMBA1000501	64		
5	HEMBA1000504	F-HEMBA1000504	65	R-HEMBA1000504	5607
	HEMBA1000505	F-HEMBA1000505	66	R-HEMBA1000505	5608
	HEMBA1000508	F-HEMBA1000508	67	R-HEMBA1000508	5609
	HEMBA1000518	F-HEMBA1000518	68	R-HEMBA1000518	5610
10	HEMBA1000519	F-HEMBA1000519	69	R-HEMBA1000519	5611
	HEMBA1000520	F-HEMBA1000520	70	R-HEMBA1000520	5612
	HEMBA1000523	F-HEMBA1000523	71.	R-HEMBA1000523	5613
	HEMBA1000531	F-HEMBA1000531	72	R-HEMBA1000531	5614
15	HEMBA1000534	F-HEMBA1000534	73		
•	HEMBA1000540	F-HEMBA1000540	74	R-HEMBA1000540	5615
	HEMBA1000542	F-HEMBA1000542,	75		
	HEMBA1000545	F-HEMBA1000545	76	R-HEMBA1000545	5616
20	HEMBA1000555	F-HEMBA1000555	77	R-nnnnnnnnnnn	5617
20	HEMBA1000557	F-HEMBA1000557	78	R-HEMBA1000557	5618
	HEMBA1000561	F-HEMBA1000561	79	R-HEMBA1000561	5619
	HEMBA1000563	F-HEMBA1000563	80	R-HEMBA1000563	5620
	HEMBA1000568	F-HEMBA1000568	81	R-HEMBA1000568	5621
25	HEMBA1000569	F-HEMBA1000569	82	R-nnnnnnnnnnn	5622
	HEMBA1000575	F-HEMBA1000575	83	R-HEMBA1000575	5623
	HEMBA1000588	F-HEMBA1000588	84	R-HEMBA1000588	5624
	HEMBA1000591	F-HEMBA1000591	85	R-HEMBA1000591	5625
30	HEMBA1000592	F-HEMBA1000592	86	R-HEMBA1000592	5626
	HEMBA1000594	F-HEMBA1000594	87	R-HEMBA1000594	5627
	HEMBA1000604	F-HEMBA1000604	88	R-HEMBA1000604	5628
	HEMBA1000608	F-HEMBA1000608	89	R-HEMBA1000608	5629
35	HEMBA1000622	F-HEMBA1000622	90	R-HEMBA1000622	5630
	HEMBA1000636	F-HEMBA1000636	91	R-HEMBA1000636	5631
	HEMBA1000637	F-HEMBA1000637	92	R-HEMBA1000637	5632
	HEMBA1000655	F-HEMBA1000655	93	R-HEMBA1000655	5633
40	HEMBA1000657	F-HEMBA1000657	94	R-HEMBA1000657	5634
	HEMBA1000662	F-HEMBA1000662	95	R-HEMBA1000662	5635
	HEMBA1000673	F-HEMBA1000673	96	R-HEMBA1000673	5636
	HEMBA1000682	F-HEMBA1000682	97	R-HEMBA1000682	5637
45	HEMBA1000686	F-HEMBA1000686	98	R-HEMBA1000686	5638
73	HEMBA1000702	F-HEMBA1000702	99	R-HEMBA1000702	5639
	HEMBA1000705	F-HEMBA1000705	100	R-HEMBA1000705	5640
	HEMBA1000719	F-HEMBA1000719	101	R-HEMBA1000719	5641
	HEMBA1000722	F-HEMBA1000722	102	R-HEMBA1000722	5642
50	HEMBA1000726	F-HEMBA1000726	103	R-HEMBA1000726	5643
	HEMBA1000727	F-HEMBA1000727	104	R-HEMBA1000727	5644
	HEMBA1000747	F-HEMBA1000747	105	R-HEMBA1000747	5645
	HEMBA1000749	F-HEMBA1000749	106	R-HEMBA1000749	5646
55	HEMBA1000752	F-HEMBA1000752	107	R-HEMBA1000752	5647

	HEMBA1000769	F-HEMBA1000769	108	R-HEMBA1000769	5648
	HEMBA1000773	F-HEMBA1000773	109	R-HEMBA1000773	5649
	HEMBA1000774	F-HEMBA1000774	110	R-HEMBA1000774	5650
5	HEMBA1000791	F-HEMBA1000791	111	R-HEMBA1000791	5651
	HEMBA1000817	F-HEMBA1000817	112	R-HEMBA1000817	5652
	HEMBA1000822	F-HEMBA1000822	113	R-HEMBA1000822	5653
	HEMBA1000827	F-HEMBA1000827	114	R-HEMBA1000827	5654
10	HEMBA1000843	F-HEMBA1000843	115	R-HEMBA1000843	5655
	HEMBA1000851	F-HEMBA1000851	116	R-HEMBA1000851	5656
	HEMBA1000852	F-HEMBA1000852	117	R-HEMBA1000852	5657
	HEMBA1000867	F-HEMBA1000867	118	R-HEMBA1000867	5658
15	HEMBA1000869	F-HEMBA1000869	119	R-HEMBA1000869	5659
15	HEMBA1000870	F-HEMBA1000870	120	R-HEMBA1000870	5660
	HEMBA1000872	F-HEMBA1000872,	121	R-HEMBA1000872	5661
	HEMBA1000876	F-HEMBA1000876	122	R-HEMBA1000876	5662
	HEMBA1000908	F-HEMBA1000908	123	R-HEMBA1000908	5663
20	HEMBA1000910	F-HEMBA1000910	124	R-HEMBA1000910	5664
	HEMBA1000918	F-HEMBA1000918	125	R-HEMBA1000918	5665
	HEMBA1000919	F-HEMBA1000919	126		5666
	HEMBA1000934	F-HEMBA1000934	127	R-HEMBA1000934	5667
25	HEMBA1000942	F-HEMBA1000942	128	R-HEMBA1000942	5668
	HEMBA1000943	F-HEMBA1000943	129	R-HEMBA1000943	5669
	HEMBA1000946	F-HEMBA1000946	130	R-HEMBA1000946	5670
	HEMBA1000960	F-HEMBA1000960	131	R-HEMBA1000960	5671
30	HEMBA1000968	F-HEMBA1000968	132	R-HEMBA1000968	5672
	HEMBA1000971	F-HEMBA1000971	133	R-HEMBA1000971	5673
	HEMBA1000972	F-HEMBA1000972	134	R-HEMBA1000972	5674
	HEMBA1000974	F-HEMBA1000974	135	R-HEMBA1000974	5675
35	HEMBA1000975	F-HEMBA1000975	136	R-HEMBA1000975	5676
	HEMBA1000985	F-HEMBA1000985	137	R-HEMBA1000985	5677
	HEMBA1000986	F-HEMBA1000986	138	R-HEMBA1000986	5678
	HEMBA1000991	F-HEMBA1000991	139	R-HEMBA1000991	5679
40	HEMBA1001007	F-HEMBA1001007	140	R-HEMBA1001007	5680
	HEMBA1001008	F-HEMBA1001008	141	R-HEMBA1001008	5681
	HEMBA1001009	F-HEMBA1001009	142	R-HEMBA1001009	5682
	HEMBA1001017	F-HEMBA1001017	143	R-HEMBA1001017	5683
45	HEMBA1001019	F-HEMBA1001019	144	R-HEMBA1001019	5684
45	HEMBA1001020	F-HEMBA1001020	145	R-HEMBA1001020	5685
	HEMBA1001022	F-HEMBA1001022	146	R-HEMBA1001022	5686
	HEMBA1001024	F-HEMBA1001024	147	R-HEMBA1001024	5687
	HEMBA1001026	F-HEMBA1001026	148	R-HEMBA1001026	5688
50	HEMBA1001043	F-HEMBA1001043	149	R-nnnnnnnnnnn	5689
	HEMBA1001051	F-HEMBA1001051	150	R-HEMBA1001051	5690
	HEMBA1001052	F-HEMBA1001052	151	R-HEMBA1001052	5691
	HEMBA1001059	F-HEMBA1001059	152	•	
55	HEMBA1001060	F-HEMBA1001060	153	R-HEMBA1001060	5692

		•			
	·HEMBA1001071	F-HEMBA1001071	154	R-HEMBA1001071	5693
	HEMBA1001077	F-HEMBA1001077	155	R-HEMBA1001077	5694
	HEMBA1001080	F-HEMBA1001080	156	R-HEMBA1001080	5695
5	HEMBA1001085	F-HEMBA1001085	157	R-HEMBA1001085	5696
	HEMBA1001088	F-HEMBA1001088	158	R-HEMBA1001088	5697
	HEMBA1001094	F-HEMBA1001094	159	R-HEMBA1001094	5698
	HEMBA1001099	F-HEMBA1001099	160	R-HEMBA1001099	5699
10	HEMBA1001109	F-HEMBA1001109	161	R-HEMBA1001109	5700
	HEMBA1001121	F-HEMBA1001121	162	R-HEMBA1001121	5701
	HEMBA1001122	F-HEMBA1001122	163	R-HEMBA1001122	5702
	HEMBA1001123	F-HEMBA1001123	164	R-HEMBA1001123	5703
15	HEMBA1001133	F-HEMBA1001133	165	R-HEMBA1001133	5704
,,,	HEMBA1001137	F-HEMBA1001137	166	R-HEMBA1001137	5705
	HEMBA1001140	F-HEMBA1001140,	167	R-HEMBA1001140	5706
	HEMBA1001172	F-HEMBA1001172	168	R-HEMBA1001172	5707
	HEMBA1001174	F-HEMBA1001174	169	R-HEMBA1001174	5708
20	HEMBA1001197	F-HEMBA1001197	170	R-HEMBA1001197	5709
	HEMBA1001208	F-HEMBA1001208	171	R-HEMBA1001208	5710
	HEMBA1001213	F-HEMBA1001213	172	•	
	HEMBA1001226	F-HEMBA1001226	173	R-HEMBA1001226	5711
25	HEMBA1001235	F-HEMBA1001235	174	R-HEMBA1001235	5712
	HEMBA1001247	F-HEMBA1001247	175	R-HEMBA1001247	5713
	HEMBA1001257	F-HEMBA1001257	176	R-HEMBA1001257	5714
	HEMBA1001265	F-HEMBA1001265	177	R-HEMBA1001265	5715
30	HEMBA1001281	F-HEMBA1001281	178	R-nnnnnnnnnnn	5716
	HEMBA1001286	F-HEMBA1001286	179	R-HEMBA1001286	5717
	HEMBA1001289	F-HEMBA1001289	180	R-HEMBA1001289	5718
	HEMBA1001294	F-HEMBA1001294	181	R-HEMBA1001294	5719
35	HEMBA1001299	F-HEMBA1001299	182	R-HEMBA1001299	5720
	HEMBA1001302	F-HEMBA1001302	183	R-HEMBA1001302	5721
	HEMBA1001303	F-HEMBA1001303	184	R-HEMBA1001303	5722
	HEMBA1001310	F-HEMBA1001310	185	R-HEMBA1001310	5723
40	HEMBA1001319	F-HEMBA1001319	186	R-HEMBA1001319	5724
	HEMBA1001323	F-HEMBA1001323	187	R-HEMBA1001323	5725
	HEMBA1001326	F-HEMBA1001326	188	R-HEMBA1001326	5726
	HEMBA1001327	F-HEMBA1001327	189	R-HEMBA1001327	5727
45	HEMBA1001330		190	R-HEMBA1001330	5728
,5	HEMBA1001351		191	R-HEMBA1001351	5729
	HEMBA1001361		192	R-HEMBA1001361	5730
	HEMBA1001375		193	R-HEMBA1001375	5731
	HEMBA1001377		194	R-HEMBA1001377	5732
50	HEMBA1001383		195	R-HEMBA1001383	5733 5734
	HEMBA1001387		196	R-HEMBA1001387	
	HEMBA1001388		197	R-HEMBA1001388	5735
	HEMBA1001391		198	R-HEMBA1001391	5736
<i>55</i>	HEMBA1001398	F-HEMBA1001398	199	R-HEMBA1001398	5737

	HEMBA1001405	F-HEMBA1001405	200	R-HEMBA1001405	5738
•	HEMBA1001407	F-HEMBA1001407	201	R-HEMBA1001407	5739
	HEMBA1001411	F-HEMBA1001411	202	R-HEMBA1001411	5740
5	HEMBA1001413	F-HEMBA1001413	203	R-HEMBA1001413	5741
	HEMBA1001415	F-HEMBA1001415	204	R-HEMBA1001415	5742
	HEMBA1001432	F-HEMBA1001432	205	R-HEMBA1001432	5743
	HEMBA1001433	F-HEMBA1001433	206	R-HEMBA1001433	5744
10	HEMBA1001435	F-HEMBA1001435	207	R-HEMBA1001435	5745
	HEMBA1001442	F-HEMBA1001442	208	R-HEMBA1001442	5746
	HEMBA1001446	F-HEMBA1001446	209	R-HEMBA1001446	5747
	HEMBA1001450	F-HEMBA1001450	210	R-HEMBA1001450	5748
15	HEMBA1001454	F-HEMBA1001454	21:1	R-HEMBA1001454	5749
13	HEMBA1001455	F-HEMBA1001455	212	R-HEMBA1001455	5750
	HEMBA1001463	F-HEMBA1001463,	213	R-HEMBA1001463	5751
	HEMBA1001476	F-HEMBA1001476	214	R-HEMBA1001476	5752
	HEMBA1001478	F-HEMBA1001478	215	R-HEMBA1001478	5753
20	HEMBA1001497	F-HEMBA1001497	216	R-HEMBA1001497	5754
	HEMBA1001510	F-HEMBA1001510	217	R-HEMBA1001510	5755
	HEMBA1001515	F-HEMBA1001515	218	R-HEMBA1001515	5756
	HEMBA1001517	F-HEMBA1001517	219	R-HEMBA1001517	5757
25	HEMBA1001522	F-HEMBA1001522	220	R-HEMBA1001522	5758
	HEMBA1001526	F-HEMBA1001526	221	R-HEMBA1001526	5759
	HEMBA1001533	F-HEMBA1001533	222	R-HEMBA1001533	5760
	HEMBA1001557	F-HEMBA1001557	223	R-HEMBA1001557	5761
30	HEMBA1001566	F-HEMBA1001566	224	R-HEMBA1001566	5762
	HEMBA1001569	F-HEMBA1001569	225	R-HEMBA1001569	5763
	HEMBA1001570	F-HEMBA1001570	226	R-HEMBA1001570	5764
	HEMBA1001579	F-HEMBA1001579	227	R-HEMBA1001579	5765
35	HEMBA1001581	F-HEMBA1001581	228	R-HEMBA1001581	5766
	HEMBA1001585	F-HEMBA1001585	229	R-HEMBA1001585	5767
	HEMBA1001589	F-HEMBA1001589	230	R-HEMBA1001589	5768
,	HEMBA1001595	F-HEMBA1001595	231	R-HEMBA1001595	5769
40	HEMBA1001608	F-HEMBA1001608	232	R-HEMBA1001608	5770
	HEMBA1001620	F-HEMBA1001620	233	R-HEMBA1001620	5771
	HEMBA1001635	F-HEMBA1001635	234	R-nannanananan	5772
	HEMBA1001636	F-HEMBA1001636	235	R-HEMBA1001636	5773
45	HEMBA1001640	F-HEMBA1001640	236	R-HEMBA1001640	5774
45	HEMBA1001647	F-HEMBA1001647	237		
	HEMBA1001651	F-HEMBA1001651	238	R-nnnnnnnnnnnn	5775
	HEMBA1001655	F-HEMBA1001655	239	R-HEMBA1001655	5776
	HEMBA1001658	F-HEMBA1001658	240	R-HEMBA1001658	5777
50	HEMBA1001661	F-HEMBA1001661	241	R-HEMBA1001661	5778
	HEMBA1001672	F-HEMBA1001672	242	R-HEMBA1001672	5779
	HEMBA1001675	F-HEMBA1001675	243	R-HEMBA1001675	5780
	HEMBA1001678	F-HEMBA1001678	244	R-HEMBA1001678	5781
55	HEMBA1001681	F-HEMBA1001681	245	R-HEMBA1001681	5782

HEMBA1001709		HEMBA1001702	F-HEMBA1001702	246	R-HEMBA1001702	5783
HEMBA1001711		HEMBA1001709				
HEMBA1001712		HEMBA1001711				
HEMBA1001714	5					
HEMBA1001718		•	. "			
HEMBA1001723						
HEMBA1001731   F-HEMBA1001731   5790     HEMBA1001734   F-HEMBA1001734   254   R-HEMBA1001734   5791     HEMBA1001745   F-HEMBA1001742   255   R-HEMBA1001745   5793     HEMBA1001746   F-HEMBA1001746   257   R-HEMBA1001746   5794     HEMBA1001746   F-HEMBA1001761   258   R-HEMBA1001761   5795     HEMBA1001781   F-HEMBA1001781   259   R-HEMBA1001781   5796     HEMBA1001781   F-HEMBA1001781   259   R-HEMBA1001781   5796     HEMBA1001791   F-HEMBA1001781   261   R-HEMBA1001781   5798     HEMBA1001800   F-HEMBA1001800   262   R-HEMBA1001800   5799     HEMBA1001800   F-HEMBA1001800   262   R-HEMBA1001803   5800     HEMBA1001804   F-HEMBA1001804   264   R-nanpanpanpanpanpanpanpanpanpanpanpanpanp						
HEMBA1001734   F-HEMBA1001734   254   R-HEMBA1001734   5791     HEMBA1001746   F-HEMBA1001745   256   R-HEMBA1001744   5792     HEMBA1001746   F-HEMBA1001745   256   R-HEMBA1001745   5793     HEMBA1001761   F-HEMBA1001761   258   R-HEMBA1001761   5795     HEMBA1001784   F-HEMBA1001781   259   R-HEMBA1001781   5796     HEMBA1001784   F-HEMBA1001784   260   R-HEMBA1001781   5796     HEMBA1001804   F-HEMBA1001803   261   R-HEMBA1001804   5799     HEMBA1001800   F-HEMBA1001803   262   R-HEMBA1001803   5800     HEMBA1001800   F-HEMBA1001803   263   R-HEMBA1001803   5800     HEMBA1001804   F-HEMBA1001804   264   R-INDIRDININA   5800     HEMBA1001805   F-HEMBA1001806   265   R-HEMBA1001808   5802     HEMBA1001806   F-HEMBA1001808   265   R-HEMBA1001808   5802     HEMBA1001809   F-HEMBA1001809   266   R-HEMBA1001808   5802     HEMBA1001815   F-HEMBA1001815   267   R-HEMBA1001815   5804     HEMBA1001820   F-HEMBA1001820   269   R-HEMBA1001819   5805     HEMBA1001820   F-HEMBA1001820   269   R-HEMBA1001820   5806     HEMBA1001824   F-HEMBA1001822   270   R-INDIRDININDIN   5801     HEMBA1001824   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001825   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001826   F-HEMBA1001824   275   R-HEMBA1001824   5808     HEMBA1001844   F-HEMBA1001847   274   R-HEMBA1001844   5810     HEMBA1001866   F-HEMBA1001867   275   R-HEMBA1001866   5814     HEMBA1001866   F-HEMBA1001867   276   R-HEMBA1001866   5814     HEMBA1001868   F-HEMBA1001869   278   R-HEMBA1001866   5814     HEMBA1001869   F-HEMBA1001869   278   R-HEMBA1001866   5814     HEMBA1001910   F-HEMBA1001910   281   R-HEMBA1001910   5818     HEMBA1001910   F-HEMBA1001911   282   R-HEMBA1001912   5819     HEMBA1001915   F-HEMBA1001913   283   R-HEMBA1001915   5820     HEMBA1001915   F-HEMBA1001915   284   R-HEMBA1001915   5820     HEMBA1001916   F-HEMBA1001916   284   R-HEMBA1001915   5820     HEMBA1001916   F-HEMBA1001916   288   R-HEMBA1001915   5820     HEMBA1001916   F-HEMBA1001916   288   R-HEMBA1001915   5820	10					
HEMBA1001744   F-HEMBA1001744   255   R-HEMBA1001745   5793						
HEMBA1001745						
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HEMBA1001761	45					
HEMBA1001781   F-HEMBA1001781   259   R-HEMBA1001781   5796     HEMBA1001784   F-HEMBA1001784   260   R-HEMBA1001784   5797     HEMBA1001791   F-HEMBA1001791   261   R-HEMBA1001791   5798     HEMBA1001800   F-HEMBA1001803   262   R-HEMBA1001800   5799     HEMBA1001804   F-HEMBA1001803   263   R-HEMBA1001803   5800     HEMBA1001805   F-HEMBA1001804   264   R-INDIRIDATION   7800     HEMBA1001806   F-HEMBA1001808   265   R-HEMBA1001808   5802     HEMBA1001815   F-HEMBA1001809   266   R-HEMBA1001809   5803     HEMBA1001815   F-HEMBA1001815   267   R-HEMBA1001815   5804     HEMBA1001819   F-HEMBA1001815   268   R-HEMBA1001819   5805     HEMBA1001820   F-HEMBA1001820   269   R-HEMBA1001820   5806     HEMBA1001824   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001824   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001824   F-HEMBA1001835   272   R-HEMBA1001824   5809     HEMBA1001824   F-HEMBA1001835   272   R-HEMBA1001824   5809     HEMBA1001824   F-HEMBA1001835   272   R-HEMBA1001824   5810     HEMBA1001844   F-HEMBA1001844   273   R-HEMBA1001847   5811     HEMBA1001866   F-HEMBA1001867   274   R-HEMBA1001847   5811     HEMBA1001866   F-HEMBA1001867   275   R-HEMBA1001847   5811     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001866   5813     HEMBA1001866   F-HEMBA1001868   276   R-HEMBA1001866   5814     HEMBA1001866   F-HEMBA1001868   278   R-HEMBA1001866   5814     HEMBA1001910   F-HEMBA1001910   281   R-HEMBA1001910   5818     HEMBA1001911   F-HEMBA1001910   281   R-HEMBA1001910   5818     HEMBA1001915   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001915   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001916   288   R-HEMBA1001915   5821     HEMBA1001917   F-HEMBA1001918   285   R-HEMBA1001915   5821     HEMBA1001917   F-HEMBA1001918   285   R-HEMBA1001915   5821     HEMBA1001917   F-HEMBA1001918   286   R-HEMBA1001915   5825     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001940   5825     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001945   5825	15		•			
HEMBA1001784   F-HEMBA1001784   260   R-HEMBA1001784   5797     HEMBA1001791   F-HEMBA1001791   261   R-HEMBA1001791   5798     HEMBA1001800   F-HEMBA1001800   262   R-HEMBA1001800   5799     HEMBA1001803   F-HEMBA1001803   263   R-HEMBA1001803   5800     HEMBA1001804   F-HEMBA1001808   265   R-HEMBA1001808   5802     HEMBA1001809   F-HEMBA1001809   266   R-HEMBA1001809   5803     HEMBA1001815   F-HEMBA1001815   267   R-HEMBA1001815   5804     HEMBA1001819   F-HEMBA1001819   268   R-HEMBA1001819   5805     HEMBA1001820   F-HEMBA1001820   269   R-HEMBA1001820   5806     HEMBA1001824   F-HEMBA1001822   270   R-nnnnnnnnnnn   5807     HEMBA1001824   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001844   F-HEMBA1001844   271   R-HEMBA1001824   5809     HEMBA1001844   F-HEMBA1001844   271   R-HEMBA1001835   5809     HEMBA1001844   F-HEMBA1001844   271   R-HEMBA1001844   5810     HEMBA1001846   F-HEMBA1001847   274   R-HEMBA1001844   5810     HEMBA1001866   F-HEMBA1001861   275   R-HEMBA1001861   5812     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001864   5813     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001864   5813     HEMBA1001866   F-HEMBA1001866   278   R-HEMBA1001864   5813     HEMBA1001866   F-HEMBA1001866   278   R-HEMBA1001866   5814     HEMBA1001886   F-HEMBA1001888   279   R-HEMBA1001889   5816     HEMBA1001910   F-HEMBA1001910   281   R-HEMBA1001910   5818     HEMBA1001915   F-HEMBA1001912   282   R-HEMBA1001912   5819     HEMBA1001915   F-HEMBA1001913   283   R-HEMBA1001913   5820     HEMBA1001916   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001916   286   R-HEMBA1001915   5821     HEMBA1001917   F-HEMBA1001919   287   R-HEMBA1001915   5821     HEMBA1001919   F-HEMBA1001919   287   R-HEMBA1001915   5821     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001940   5825     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001942   5826     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001942   5826     HEMBA1001940   F-HEMBA1001945   290   R-HEMBA1001945   5826						
HEMBA1001801   F-HEMBA1001791   261   R-HEMBA1001791   5798     HEMBA1001800   F-HEMBA1001800   262   R-HEMBA1001800   5799     HEMBA1001803   F-HEMBA1001803   263   R-HEMBA1001803   5800     HEMBA1001804   F-HEMBA1001804   264   R-INDIRDININDININDININDININDININDININDIN			•			
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HEMBA1001804   F-HEMBA1001804   264   R-nnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnn						
HEMBA1001808   F-HEMBA1001808   265   R-HEMBA1001808   5802     HEMBA1001809   F-HEMBA1001809   266   R-HEMBA1001809   5803     HEMBA1001815   F-HEMBA1001815   267   R-HEMBA1001815   5804     HEMBA1001819   F-HEMBA1001819   268   R-HEMBA1001819   5805     HEMBA1001820   F-HEMBA1001820   269   R-HEMBA1001820   5806     HEMBA1001822   F-HEMBA1001822   270   R-nnnnnnnnnnn   5807     HEMBA1001835   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001835   F-HEMBA1001835   272   R-HEMBA1001835   5809     HEMBA1001844   F-HEMBA1001844   273   R-HEMBA1001844   5810     HEMBA1001847   F-HEMBA1001847   274   R-HEMBA1001847   5811     HEMBA1001861   F-HEMBA1001861   275   R-HEMBA1001861   5812     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001864   5813     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001866   5814     HEMBA1001869   F-HEMBA1001888   279   R-HEMBA1001866   5814     HEMBA1001910   F-HEMBA1001910   281   R-HEMBA1001910   5818     HEMBA1001911   F-HEMBA1001912   282   R-HEMBA1001910   5818     HEMBA1001915   F-HEMBA1001911   281   R-HEMBA1001910   5818     HEMBA1001915   F-HEMBA1001912   282   R-HEMBA1001912   5820     HEMBA1001916   F-HEMBA1001913   283   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001917   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001918   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001915   286   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001915   286   R-HEMBA1001915   5823     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001940   5825     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001940   5825						
HEMBA1001809						
HEMBA1001815    F-HEMBA1001815    267    R-HEMBA1001815    5804      HEMBA1001819    F-HEMBA1001819    268    R-HEMBA1001819    5805      HEMBA1001820    F-HEMBA1001820    269    R-HEMBA1001820    5806      HEMBA1001822    F-HEMBA1001822    270    R-nnnnnnnnnnn	25					
HEMBA1001819 F-HEMBA1001819 268 R-HEMBA1001819 5805 HEMBA1001820 F-HEMBA1001820 269 R-HEMBA1001820 5806 HEMBA1001822 F-HEMBA1001822 270 R-nnnnnnnnnnn 5807 HEMBA1001824 F-HEMBA1001824 271 R-HEMBA1001824 5808 HEMBA1001835 F-HEMBA1001835 272 R-HEMBA1001835 5809 HEMBA1001844 F-HEMBA1001844 273 R-HEMBA1001844 5810 HEMBA1001847 F-HEMBA1001847 274 R-HEMBA1001847 5811 HEMBA1001861 F-HEMBA1001861 275 R-HEMBA1001861 5812 HEMBA1001864 F-HEMBA1001864 276 R-HEMBA1001864 5813 HEMBA1001866 F-HEMBA1001866 277 R-HEMBA1001866 5814 HEMBA1001888 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001914 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001919 F-HEMBA1001912 286 R-HEMBA1001913 5822 HEMBA1001919 F-HEMBA1001912 286 R-HEMBA100191 5823 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001942 289 R-HEMBA1001942 5826						
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HEMBA1001822   F-HEMBA1001822   270   R-nnnnnnnnnnn   5807     HEMBA1001824   F-HEMBA1001824   271   R-HEMBA1001824   5808     HEMBA1001835   F-HEMBA1001835   272   R-HEMBA1001835   5809     HEMBA1001844   F-HEMBA1001844   273   R-HEMBA1001844   5810     HEMBA1001847   F-HEMBA1001847   274   R-HEMBA1001847   5811     HEMBA1001861   F-HEMBA1001861   275   R-HEMBA1001861   5812     HEMBA1001864   F-HEMBA1001864   276   R-HEMBA1001864   5813     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001866   5814     HEMBA1001869   F-HEMBA1001869   278   R-nnnnnnnnnnn   5815     HEMBA1001888   F-HEMBA1001888   279   R-HEMBA1001888   5816     HEMBA1001896   F-HEMBA1001888   279   R-HEMBA1001886   5817     HEMBA1001910   F-HEMBA1001910   281   R-HEMBA1001910   5818     HEMBA1001912   F-HEMBA1001912   282   R-HEMBA1001912   5819     HEMBA1001913   F-HEMBA1001913   283   R-HEMBA1001913   5820     HEMBA1001915   F-HEMBA1001915   284   R-HEMBA1001915   5821     HEMBA1001916   F-HEMBA1001918   285   R-HEMBA1001915   5821     HEMBA1001917   F-HEMBA1001918   285   R-HEMBA1001915   5823     HEMBA1001921   F-HEMBA1001921   286   R-HEMBA1001921   5823     HEMBA1001939   F-HEMBA1001940   288   R-HEMBA1001940   5825     HEMBA1001940   F-HEMBA1001940   288   R-HEMBA1001940   5825     HEMBA1001940   F-HEMBA1001942   289   R-HEMBA1001942   5826     HEMBA1001945   F-HEMBA1001945   290   R-HEMBA1001945   5827						
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HEMBA1001835 F-HEMBA1001835 272 R-HEMBA1001835 5809 HEMBA1001844 F-HEMBA1001844 273 R-HEMBA1001844 5810 HEMBA1001847 F-HEMBA1001847 274 R-HEMBA1001847 5811 HEMBA1001861 F-HEMBA1001861 275 R-HEMBA1001861 5812 HEMBA1001864 F-HEMBA1001864 276 R-HEMBA1001864 5813 HEMBA1001866 F-HEMBA1001866 277 R-HEMBA1001866 5814 HEMBA1001869 F-HEMBA1001869 278 R-nnnnnnnnnnnn 5815 HEMBA1001888 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001916 F-HEMBA1001912 285 R-HEMBA1001915 5821 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827					R-HEMBA1001824	
HEMBA1001847 F-HEMBA1001847 274 R-HEMBA1001847 5811 HEMBA1001861 F-HEMBA1001861 275 R-HEMBA1001861 5812 HEMBA1001864 F-HEMBA1001864 276 R-HEMBA1001864 5813 HEMBA1001866 F-HEMBA1001866 277 R-HEMBA1001866 5814 HEMBA1001889 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001919 F-HEMBA1001915 286 R-HEMBA1001918 5822 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5826						
HEMBA1001847   F-HEMBA1001847   274   R-HEMBA1001847   5811     HEMBA1001861   F-HEMBA1001861   275   R-HEMBA1001861   5812     HEMBA1001864   F-HEMBA1001864   276   R-HEMBA1001864   5813     HEMBA1001866   F-HEMBA1001866   277   R-HEMBA1001866   5814     HEMBA1001869   F-HEMBA1001869   278   R-INININININININININININININININININININ		HEMBA1001844	F-HEMBA1001844		R-HEMBA1001844	
HEMBA1001861 F-HEMBA1001861 275 R-HEMBA1001861 5812 HEMBA1001864 F-HEMBA1001864 276 R-HEMBA1001864 5813 HEMBA1001866 F-HEMBA1001866 277 R-HEMBA1001866 5814 HEMBA1001889 F-HEMBA1001889 278 R-nnnnnnnnnnn 5815 HEMBA1001896 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827	35	HEMBA1001847	F-HEMBA1001847		R-HEMBA1001847	5811
HEMBA1001866 F-HEMBA1001866 277 R-HEMBA1001866 5814 HEMBA1001869 F-HEMBA1001869 278 R-nnnnnnnnnnn 5815 HEMBA1001888 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001861	F-HEMBA1001861		R-HEMBA1001861	5812
HEMBA1001869 F-HEMBA1001869 278 R-nnnnnnnnnnn 5815 HEMBA1001888 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001864	F-HEMBA1001864	276	R-HEMBA1001864	5813
HEMBA1001888 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001866	F-HEMBA1001866	277	R-HEMBA1001866	5814
HEMBA1001888 F-HEMBA1001888 279 R-HEMBA1001888 5816 HEMBA1001896 F-HEMBA1001896 280 R-HEMBA1001896 5817 HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827	40	HEMBA1001869	F-HEMBA1001869	278	R-nnnnnnnnnnn	5815
HEMBA1001910 F-HEMBA1001910 281 R-HEMBA1001910 5818 HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827	,	HEMBA1001888	F-HEMBA1001888	279	R-HEMBA1001888	5816
HEMBA1001912 F-HEMBA1001912 282 R-HEMBA1001912 5819 HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001896	F-HEMBA1001896	280	R-HEMBA1001896	5817
HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001910	F-HEMBA1001910	281	R-HEMBA1001910	5818
HEMBA1001913 F-HEMBA1001913 283 R-HEMBA1001913 5820 HEMBA1001915 F-HEMBA1001915 284 R-HEMBA1001915 5821 HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827	45	HEMBA1001912	F-HEMBA1001912	282	R-HEMBA1001912	5819
HEMBA1001918 F-HEMBA1001918 285 R-HEMBA1001918 5822 HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823 HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824 HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827	45	HEMBA1001913	F-HEMBA1001913	283	R-HEMBA1001913	5820
HEMBA1001921 F-HEMBA1001921 286 R-HEMBA1001921 5823  HEMBA1001939 F-HEMBA1001939 287 R-HEMBA1001939 5824  HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825  HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826  HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001915	F-HEMBA1001915	284	R-HEMBA1001915	5821
50         HEMBA1001939         F-HEMBA1001939         287         R-HEMBA1001939         5824           HEMBA1001940         F-HEMBA1001940         288         R-HEMBA1001940         5825           HEMBA1001942         F-HEMBA1001942         289         R-HEMBA1001942         5826           HEMBA1001945         F-HEMBA1001945         290         R-HEMBA1001945         5827		HEMBA1001918	F-HEMBA1001918	285	R-HEMBA1001918	5822
HEMBA1001940 F-HEMBA1001940 288 R-HEMBA1001940 5825 HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001921	F-HEMBA1001921	286	R-HEMBA1001921	5823
HEMBA1001942 F-HEMBA1001942 289 R-HEMBA1001942 5826 HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827	50	HEMBA1001939	F-HEMBA1001939	287	R-HEMBA1001939	5824
HEMBA1001945 F-HEMBA1001945 290 R-HEMBA1001945 5827		HEMBA1001940	F-HEMBA1001940	288	R-HEMBA1001940	5825
		HEMBA1001942	F-HEMBA1001942	289	R-HEMBA1001942	5826
55 HEMBA1001950 F-HEMBA1001950 291 R-HEMBA1001950 5828		HEMBA1001945	F-HEMBA1001945	290	R-HEMBA1001945	5827
	55	HEMBA1001950	F-HEMBA1001950	291	R-HEMBA1001950	5828

	HEMBA1001960	F-HEMBA1001960	292	R-HEMBA1001960	5829
	HEMBA1001962	F-HEMBA1001962	293	R-HEMBA1001962	5830
	HEMBA1001964	F-HEMBA1001964	294	R-HEMBA1001964	5831
5 ,	HEMBA1001967	F-HEMBA1001967	295	R-HEMBA1001967	5832
	HEMBA1001979	F-HEMBA1001979	296	R-HEMBA1001979	5833
	HEMBA1001987	F-HEMBA1001987	297	R-HEMBA1001987	5834
	HEMBA1001991	F-HEMBA1001991	298	R-HEMBA1001991	5835
10	HEMBA1002003	F-HEMBA1002003	299	R-HEMBA1002003	5836
	HEMBA1002008	F-HEMBA1002008	300	R-HEMBA1002008	5837
	HEMBA1002018	F-HEMBA1002018	301	R-HEMBA1002018	5838
	HEMBA1002022	F-HEMBA1002022	302	R-HEMBA1002022	5839
	HEMBA1002035	F-HEMBA1002035	303	R-HEMBA1002035	5840
15	HEMBA1002039	F-HEMBA1002039	304	R-HEMBA1002039	5841
	HEMBA1002049	F-HEMBA1002049	305	R-HEMBA1002049	5842
	HEMBA1002084	F-HEMBA1002084	306	R-HEMBA1002084	5843
	HEMBA1002092	F-HEMBA1002092	307	R-HEMBA1002092	5844
20	HEMBA1002100	F-HEMBA1002100	308	R-HEMBA1002100	5845
	HEMBA1002102	F-HEMBA1002102	309	R-HEMBA1002102	5846
	HEMBA1002113	F-HEMBA1002113	310	R-HEMBA1002113	5847
	HEMBA1002119	F-HEMBA1002119	311	R-HEMBA1002119	5848
25	HEMBA1002125	F-HEMBA1002125	312	R-HEMBA1002125	5849
	HEMBA1002139	F-HEMBA1002139	313	R-HEMBA1002139	5850
	HEMBA1002144	F-HEMBA1002144	314	R-HEMBA1002144	5851
	HEMBA1002150	F-HEMBA1002150	315	R-HEMBA1002150	5852
30	HEMBA1002151	F-HEMBA1002151	316	R-HEMBA1002151	5853
	HEMBA1002153	F-HEMBA1002153	317	R-HEMBA1002153	5854
	HEMBA1002160	F-HEMBA1002160	318	R-HEMBA1002160	5855
	HEMBA1002161	F-HEMBA1002161	319	R-HEMBA1002161	5856
35	HEMBA1002162	F-HEMBA1002162	320	R-HEMBA1002162	5857
	HEMBA1002166	F-HEMBA1002166	321	R-HEMBA1002166	5858
	HEMBA1002177	F-HEMBA1002177	322	R-HEMBA1002177	5859
	HEMBA1002185	F-HEMBA1002185	323	R-HEMBA1002185	5860
	HEMBA1002189	F-HEMBA1002189	324	R-HEMBA1002189	5861
40	HEMBA1002191	F-HEMBA1002191	325	R-HEMBA1002191	5862
	HEMBA1002199	F-HEMBA1002199	326	R-HEMBA1002199	5863
	HEMBA1002204	F-HEMBA1002204	327	R-HEMBA1002204	5864
	HEMBA1002212	F-HEMBA1002212	328	R-HEMBA1002212	5865
45	HEMBA1002215	F-HEMBA1002215	329	R-HEMBA1002215	5866
	HEMBA1002226	F-HEMBA1002226	330	R-HEMBA1002226	5867
	HEMBA1002229	F-HEMBA1002229	331	R-HEMBA1002229	5868
	HEMBA1002237	F-HEMBA1002237	332	R-HEMBA1002237	5869
50	HEMBA1002241	F-HEMBA1002241	333		
	HEMBA1002253	F-HEMBA1002253	334	R-HEMBA1002253	5870
	HEMBA1002257	F-HEMBA1002257	335	R-HEMBA1002257	5871
	HEMBA1002265	F-HEMBA1002265	336		
55	HEMBA1002267	F-HEMBA1002267	337	R-HEMBA1002267	5872

HEMBA1002321   F-HEMBA1002321   339   R-HEMBA1002321   536     HEMBA1002337   F-HEMBA1002337   341   R-HEMBA1002337   536     HEMBA1002341   F-HEMBA1002341   342   R-HEMBA1002343   536     HEMBA1002342   F-HEMBA1002343   342   R-HEMBA1002343   536     HEMBA1002343   F-HEMBA1002344   342   R-HEMBA1002349   536     HEMBA1002349   F-HEMBA1002349   344   R-HEMBA1002349   536     HEMBA1002363   F-HEMBA1002363   345   R-nannannannan   586     HEMBA1002361   F-HEMBA1002383   346   R-HEMBA1002381   346     HEMBA1002417   F-HEMBA1002313   346   R-HEMBA1002381   348     HEMBA1002417   F-HEMBA1002417   348   R-HEMBA1002381   348     HEMBA1002419   F-HEMBA1002419   349   R-HEMBA1002417   538     HEMBA1002430   F-HEMBA1002439   347   R-HEMBA1002430   536     HEMBA1002430   F-HEMBA1002439   347   R-HEMBA1002430   536     HEMBA1002430   F-HEMBA1002439   351   R-HEMBA1002430   536     HEMBA1002458   F-HEMBA1002449   357   R-HEMBA1002430   538     HEMBA1002460   F-HEMBA1002460   353   R-HEMBA1002460   538     HEMBA1002460   F-HEMBA1002460   353   R-HEMBA1002460   538     HEMBA1002460   F-HEMBA1002460   355   R-HEMBA1002460   538     HEMBA1002460   F-HEMBA1002460   358   R-HEMBA1002460   538     HEMBA1002560   F-HEMBA1002460   358   R-HEMBA1002460   538     HEMBA1002560   F-HEMBA1002560   360   R-HEMBA1002460   538     HEMBA1002560   F-HEMBA1002560   360   R-HEMBA1002560   539     HEMBA1002560   F-HEMBA1002560   370   R-HEMBA1002560   539     HEMBA1002560   F-HEMBA1002560   370   R-HEMBA1002560   539     HEMBA1002560   F-HEMBA1002560   370   R-HEMBA1002560   539     HEMBA1002560   F-HEMBA100256						
HEMBA1002328   F-HEMBA1002328   340   R-HEMBA1002328   587		HEMBA1002270.	F-HEMBA1002270	338	R-HEMBA1002270	5873
S		HEMBA1002321	F-HEMBA1002321	339	R-HEMBA1002321	5874
HEMBA1002341		HEMBA1002328 ·	F-HEMBA1002328	340	R-HEMBA1002328	5875
HEMBA1002341	5	HEMBA1002337	F-HEMBA1002337	341	R-HEMBA1002337	5876
HEMBA1002348			F-HEMBA1002341	342	R-HEMBA1002341	5877
HEMBA1002349					R-HEMBA1002348	5878
HEMBA1002363					R-HEMBA1002349	5879
HEMBA1002381	10		F-HEMBA1002363		R-nnnnnnnnnnn	5880
HEMBA1002439			F-HEMBA1002381			5881
HEMBA1002417					R-HEMBA1002389	5882
HEMBA1002430			F-HEMBA1002417	348	R-HEMBA1002417	5883
HEMBA1002430	15				R-HEMBA1002419	5884
HEMBA1002439	15				R-HEMBA1002430	5885
HEMBA1002458   F-HEMBA1002458   352   R-HEMBA1002458   588     HEMBA1002460   F-HEMBA1002460   353   R-HEMBA1002460   588     HEMBA1002462   F-HEMBA1002462   354   R-HEMBA1002462   588     HEMBA1002475   F-HEMBA1002475   356   R-nnnnnnnnnnn   588     HEMBA1002477   F-HEMBA1002477   357   R-HEMBA1002477   588     HEMBA1002486   F-HEMBA1002477   357   R-HEMBA1002477   588     HEMBA1002495   F-HEMBA1002496   358   R-HEMBA1002496   588     HEMBA1002495   F-HEMBA1002496   359   R-HEMBA1002496   588     HEMBA1002503   F-HEMBA1002498   360   R-HEMBA1002496   588     HEMBA1002503   F-HEMBA1002503   361   R-HEMBA1002503   588     HEMBA1002504   F-HEMBA1002503   362   R-HEMBA1002503   588     HEMBA1002513   F-HEMBA1002513   363   R-INNINININININININININININININININININI					R-HEMBA1002439	5886
HEMBA1002460   F-HEMBA1002460   353   R-HEMBA1002460   588     HEMBA1002462   F-HEMBA1002462   354   R-HEMBA1002462   588     HEMBA1002475   F-HEMBA1002475   356   R-Inninininininininininininininininininin			•		R-HEMBA1002458	5887
HEMBA1002462					R-HEMBA1002460	5888
HEMBA1002469   F-HEMBA1002469   355     HEMBA1002475   F-HEMBA1002475   356   R-nnnnnnnnnnn   588     HEMBA1002477   F-HEMBA1002477   357   R-HEMBA1002477   589     HEMBA1002496   F-HEMBA1002495   359   R-HEMBA1002495   589     HEMBA1002497   F-HEMBA1002495   359   R-HEMBA1002495   589     HEMBA1002503   F-HEMBA1002503   361   R-HEMBA1002503   589     HEMBA1002503   F-HEMBA1002503   361   R-HEMBA1002503   589     HEMBA1002503   F-HEMBA1002503   362   R-HEMBA1002503   589     HEMBA1002513   F-HEMBA1002513   363   R-nnnnnnnnnnnn   589     HEMBA1002513   F-HEMBA1002513   363   R-nnnnnnnnnnnn   589     HEMBA1002514   F-HEMBA1002515   364   R-HEMBA1002515   589     HEMBA1002542   F-HEMBA1002542   366   R-HEMBA1002542   599     HEMBA1002547   F-HEMBA1002542   366   R-HEMBA1002544   599     HEMBA1002555   F-HEMBA1002552   368   R-HEMBA1002554   599     HEMBA1002555   F-HEMBA1002555   369   R-HEMBA1002555   599     HEMBA1002566   F-HEMBA1002561   371   R-HEMBA1002556   599     HEMBA1002569   F-HEMBA1002569   372   R-HEMBA1002561   599     HEMBA1002569   F-HEMBA1002569   372   R-HEMBA1002561   599     HEMBA1002569   F-HEMBA1002569   374   R-HEMBA1002569   599     HEMBA1002621   F-HEMBA1002624   378   R-HEMBA1002629   599     HEMBA1002624   F-HEMBA1002624   378   R-HEMBA1002624   599     HEMBA1002626   F-HEMBA1002624   378   R-HEMBA1002625   599     HEMBA1002626   F-HEMBA1002626   379   R-HEMBA1002626   599     HEMBA1002626   F-HEMBA1002626   380   R-HEMBA1002626   599     HEMBA1002665   F-HEMBA1002665   381   R-HEMBA1002665   599     HEMBA1002665   F-HEMBA1002665   381   R-HEMBA1002665   599     HEMBA1002665   F-HEMBA1002665   382   R-HEMBA1002665   599     HEMBA1002665   F-HEMBA1002665   382   R-HEMBA1002665   599     HEMBA1002665   F-HEMBA1002665   382   R-HEMBA1002665   599     HEMBA1002665   F-HEMBA1002665   383   R-HEMBA1002665	20				R-HEMBA1002462	5889
HEMBA1002477   F-HEMBA1002477   357   R-HEMBA1002477   58		HEMBA1002469	F-HEMBA1002469	355		
### 1002486   F-HEMBA1002486   358   R-HEMBA1002486   588		HEMBA1002475	F-HEMBA1002475	356	R-nnnnnnnnnnn	5890
HEMBA1002495 F-HEMBA1002495 359 R-HEMBA1002495 58 HEMBA1002498 F-HEMBA1002498 360 R-HEMBA1002498 58 HEMBA1002503 F-HEMBA1002503 361 R-HEMBA1002503 58 HEMBA1002508 F-HEMBA1002508 362 R-HEMBA1002508 58 HEMBA1002513 F-HEMBA1002513 363 R-nnnnnnnnnnn 58 HEMBA1002515 F-HEMBA1002515 364 R-HEMBA1002515 58 HEMBA1002538 F-HEMBA1002515 364 R-HEMBA1002515 58 HEMBA1002538 F-HEMBA1002538 365 R-HEMBA1002538 58 HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002555 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002556 F-HEMBA1002558 370 R-HEMBA1002555 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002551 59 HEMBA1002569 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002583 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002591 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002590 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002609 F-HEMBA1002601 377 R-HEMBA1002602 59 HEMBA1002609 F-HEMBA1002603 379 R-HEMBA1002603 59 HEMBA1002609 F-HEMBA1002603 379 R-HEMBA1002603 59 HEMBA1002609 F-HEMBA1002603 370 R-HEMBA1002603 59 HEMBA1002609 F-HEMBA1002604 370 R-HEMBA1002605 59 HEMBA1002609 F-HEMBA1002605 380 R-HEMBA1002605 59 HEMBA1002609 F-HEMBA1002605 380 R-HEMBA1002605 59 HEMBA1002609 F-HEMBA1002605 380 R-HEMBA1002605 59		HEMBA1002477	F-HEMBA1002477	357	R-HEMBA1002477	5891
HEMBA1002498   F-HEMBA1002498   360   R-HEMBA1002498   58     HEMBA1002503   F-HEMBA1002503   361   R-HEMBA1002503   58     HEMBA1002508   F-HEMBA1002508   362   R-HEMBA1002508   58     HEMBA1002513   F-HEMBA1002513   363   R-nnnnnnnnnnn   58     HEMBA1002515   F-HEMBA1002515   364   R-HEMBA1002515   58     HEMBA1002538   F-HEMBA1002538   365   R-HEMBA1002538   58     HEMBA1002542   F-HEMBA1002542   366   R-HEMBA1002542   59     HEMBA1002547   F-HEMBA1002547   367   R-HEMBA1002547   59     HEMBA1002552   F-HEMBA1002552   368   R-HEMBA1002552   59     HEMBA1002555   F-HEMBA1002555   369   R-HEMBA1002555   59     HEMBA1002556   F-HEMBA1002558   370   R-HEMBA1002555   59     HEMBA1002561   F-HEMBA1002561   371   R-HEMBA1002561   59     HEMBA1002569   F-HEMBA1002569   372   R-nnnnnnnnnnn   59     HEMBA1002590   F-HEMBA1002590   374   R-HEMBA1002593   59     HEMBA1002590   F-HEMBA1002590   374   R-HEMBA1002592   59     HEMBA1002591   F-HEMBA1002592   375   R-HEMBA1002592   59     HEMBA1002609   F-HEMBA1002609   376     HEMBA1002609   F-HEMBA1002609   376     HEMBA1002624   F-HEMBA1002624   378   R-HEMBA1002624   59     HEMBA1002625   F-HEMBA1002624   378   R-HEMBA1002624   59     HEMBA1002626   F-HEMBA1002628   379   R-HEMBA1002624   59     HEMBA1002626   F-HEMBA1002629   380   R-HEMBA1002625   59     HEMBA100261   F-HEMBA1002629   380   R-HEMBA1002626   59     HEMBA1002651   F-HEMBA1002651   381   R-HEMBA1002625   59     HEMBA1002651   F-HEMBA1002651   382   R-HEMBA1002655   59     HEMBA1002656   F-HEMBA1002651   382   R-HEMBA1002655   59     HEMBA1002650   F-HEMBA1002651   382   R-HEMBA	25	HEMBA1002486	F-HEMBA1002486	358	R-HEMBA1002486	5892
HEMBA1002503 F-HEMBA1002503 361 R-HEMBA1002503 58 HEMBA1002508 F-HEMBA1002508 362 R-HEMBA1002508 58 HEMBA1002513 F-HEMBA1002513 363 R-nnnnnnnnnnn 58 HEMBA1002515 F-HEMBA1002515 364 R-HEMBA1002515 58 HEMBA1002538 F-HEMBA1002538 365 R-HEMBA1002538 58 HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002555 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002553 369 R-HEMBA1002555 59 HEMBA1002556 F-HEMBA1002553 370 R-HEMBA1002555 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002558 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002569 F-HEMBA1002569 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002590 F-HEMBA1002590 376 R-HEMBA1002590 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002624 F-HEMBA1002621 377 R-HEMBA1002592 59 HEMBA1002625 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002626 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002628 F-HEMBA1002623 379 R-HEMBA1002624 59 HEMBA1002629 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002626 F-HEMBA1002629 380 R-HEMBA1002625 59 HEMBA1002626 F-HEMBA1002629 380 R-HEMBA1002625 59 HEMBA1002651 F-HEMBA1002651 381 R-HEMBA1002625 59 HEMBA1002655 F-HEMBA1002659 381 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002659 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002659 383 R-HEMBA1002651 59		HEMBA1002495	F-HEMBA1002495	359	R-HEMBA1002495	5893
HEMBA1002508 F-HEMBA1002508 362 R-HEMBA1002508 58 HEMBA1002513 F-HEMBA1002513 363 R-nnnnnnnnnnn 58 HEMBA1002515 F-HEMBA1002515 364 R-HEMBA1002515 58 HEMBA1002538 F-HEMBA1002538 365 R-HEMBA1002538 58 HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002553 369 R-HEMBA1002555 59 HEMBA1002561 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002558 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002538 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002609 F-HEMBA1002690 376 HEMBA1002624 F-HEMBA1002629 375 R-HEMBA1002592 59 HEMBA1002629 F-HEMBA1002629 377 R-HEMBA1002592 59 HEMBA1002629 F-HEMBA1002629 378 R-HEMBA1002621 59 HEMBA1002629 F-HEMBA1002629 378 R-HEMBA1002621 59 HEMBA1002629 F-HEMBA1002629 378 R-HEMBA1002621 59 HEMBA1002629 F-HEMBA1002629 378 R-HEMBA1002624 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002624 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002624 59 HEMBA1002651 F-HEMBA1002651 381 R-HEMBA1002625 59 HEMBA1002655 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002659 383 R-HEMBA1002651 59		HEMBA1002498	F-HEMBA1002498	360	R-HEMBA1002498	5894
HEMBA1002513 F-HEMBA1002513 363 R-nnnnnnnnnn 58 HEMBA1002515 F-HEMBA1002515 364 R-HEMBA1002515 58 HEMBA1002538 F-HEMBA1002538 365 R-HEMBA1002538 58 HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002558 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002629 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002592 59 HEMBA1002624 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002625 F-HEMBA1002624 378 R-HEMBA1002621 59 HEMBA1002626 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002626 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002626 F-HEMBA1002628 379 R-HEMBA1002624 59 HEMBA1002626 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002626 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002626 F-HEMBA1002629 380 R-HEMBA1002625 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002656 F-HEMBA1002651 382 R-HEMBA1002655 59		HEMBA1002503	F-HEMBA1002503	361	R-HEMBA1002503	5895
HEMBA1002515 F-HEMBA1002515 364 R-HEMBA1002515 58 HEMBA1002538 F-HEMBA1002538 365 R-HEMBA1002538 58 HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002558 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002569 372 R-nnnnnnnnnnnn 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002583 59 HEMBA1002592 F-HEMBA1002590 376 R-HEMBA1002590 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002621 59 HEMBA1002625 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002626 F-HEMBA1002625 380 R-HEMBA1002625 59 HEMBA1002626 F-HEMBA1002625 381 R-HEMBA1002626 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002650 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002650 F-HEMBA1002651 382 R-HEMBA1002651 59	30	HEMBA1002508	F-HEMBA1002508	362	R-HEMBA1002508	5896
HEMBA1002538 F-HEMBA1002538 365 R-HEMBA1002538 58 HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002561 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002569 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002569 372 R-nnnnnnnnnnnn 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002583 59 HEMBA1002591 F-HEMBA1002590 375 R-HEMBA1002590 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002621 59 HEMBA1002625 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002626 F-HEMBA1002625 380 R-HEMBA1002626 59 HEMBA1002626 F-HEMBA1002629 380 R-HEMBA1002626 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002650 F-HEMBA1002651 382 R-HEMBA1002651 59		HEMBA1002513	F-HEMBA1002513	363	R-nnnnnnnnnnn	5897
HEMBA1002542 F-HEMBA1002542 366 R-HEMBA1002542 59 HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002568 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002624 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002629 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002651 F-HEMBA1002625 380 R-HEMBA1002625 59 HEMBA1002651 F-HEMBA1002651 381 R-HEMBA1002655 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002650 F-HEMBA1002651 382 R-HEMBA1002655 59		HEMBA1002515	F-HEMBA1002515	364	R-HEMBA1002515	5898
HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002558 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002624 59 HEMBA1002629 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002645 F-HEMBA1002625 381 R-HEMBA1002625 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002651 59	•	HEMBA1002538	F-HEMBA1002538	365	R-HEMBA1002538	5899
HEMBA1002547 F-HEMBA1002547 367 R-HEMBA1002547 59 HEMBA1002552 F-HEMBA1002552 368 R-HEMBA1002552 59 HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002558 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002624 59 HEMBA1002629 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002645 F-HEMBA1002625 381 R-HEMBA1002625 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002659 383 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002651 59	35	HEMBA1002542	F-HEMBA1002542	366	R-HEMBA1002542	5900
HEMBA1002555 F-HEMBA1002555 369 R-HEMBA1002555 59 HEMBA1002558 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002624 59 HEMBA1002645 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002659 F-HEMBA1002659 380 R-HEMBA1002625 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002651 F-HEMBA1002659 383 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002651 59		HEMBA1002547	F-HEMBA1002547	367	R-HEMBA1002547	5901
HEMBA1002558 F-HEMBA1002558 370 R-HEMBA1002558 59 HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002624 59 HEMBA1002645 F-HEMBA1002628 380 R-HEMBA1002629 59 HEMBA1002651 F-HEMBA1002651 381 R-HEMBA1002655 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002651 59		HEMBA1002552	F-HEMBA1002552	368		5902
HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnn		HEMBA1002555	F-HEMBA1002555	369	R-HEMBA1002555	5903
HEMBA1002561 F-HEMBA1002561 371 R-HEMBA1002561 59 HEMBA1002569 F-HEMBA1002569 372 R-nnnnnnnnnnn 59 HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002624 59 HEMBA1002645 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002650 F-HEMBA1002659 383 R-HEMBA1002651 59	40	HEMBA1002558	F-HEMBA1002558	370		5904
HEMBA1002583 F-HEMBA1002583 373 R-HEMBA1002583 59 HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002645 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002651 59	40	HEMBA1002561			R-HEMBA1002561	5905
HEMBA1002590 F-HEMBA1002590 374 R-HEMBA1002590 59 HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002650 F-HEMBA1002659 383 R-HEMBA1002659 59		HEMBA1002569	F-HEMBA1002569			5906
HEMBA1002592 F-HEMBA1002592 375 R-HEMBA1002592 59 HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59		HEMBA1002583	F-HEMBA1002583	373		5907
HEMBA1002609 F-HEMBA1002609 376 HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59		HEMBA1002590				5908
HEMBA1002621 F-HEMBA1002621 377 R-HEMBA1002621 59 HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59	45	HEMBA1002592			R-HEMBA1002592	5909
HEMBA1002624 F-HEMBA1002624 378 R-HEMBA1002624 59  HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59  HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59  HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59  HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59  HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59		HEMBA1002609				200
50 HEMBA1002628 F-HEMBA1002628 379 R-HEMBA1002628 59 HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59						5910
HEMBA1002629 F-HEMBA1002629 380 R-HEMBA1002629 59 HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59						5911
HEMBA1002645 F-HEMBA1002645 381 R-HEMBA1002645 59 HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59	50					5912
HEMBA1002651 F-HEMBA1002651 382 R-HEMBA1002651 59						5913
HEMBA1002650 F-HEMBA1002659 383 R-HEMBA1002659 59						5914
55 HEMBA1002659 F-HEMBA1002659 383 R-HEMBA1002659 59						5915
	55	HEMBA1002659	F-HEMBA1002659	383	K-HEMBA1002659	5916

	HEMBA1002661	F-HEMBA1002661	384	R-HEMBA1002661	5917
	HEMBA1002666	F-HEMBA1002666	385	R-HEMBA1002666	5918
	HEMBA1002678	F-HEMBA1002678	386	R-HEMBA1002678	5919
5	HEMBA1002679	F-HEMBA1002679	387	R-nnnnnnnnnnn	5920
	HEMBA1002688	F-HEMBA1002688	388	R-HEMBA1002688	5921
	HEMBA1002696	F-HEMBA1002696	389	R-HEMBA1002696	5922
	HEMBA1002703	F-HEMBA1002703	390		
10	HEMBA1002712	F-HEMBA1002712	391	R-HEMBA1002712	5923
	HEMBA1002716	F-HEMBA1002716	392	R-HEMBA1002716	5924
	HEMBA1002728	F-HEMBA1002728	393	R-HEMBA1002728	5925
	HEMBA1002730	F-HEMBA1002730	394	R-HEMBA1002730	5926
v.	HEMBA1002742	F-HEMBA1002742	395	R-HEMBA1002742	5927
15	HEMBA1002746	F-HEMBA1002746	396	R-HEMBA1002746	5928
	HEMBA1002748	F-HEMBA1002748,	397	R-HEMBA1002748	5929
	HEMBA1002750	F-HEMBA1002750	398	R-HEMBA1002750	5930
	HEMBA1002768	F-HEMBA1002768	399	R-HEMBA1002768	5931
20	HEMBA1002770	F-HEMBA1002770	400	R-HEMBA1002770	5932
	HEMBA1002777	F-HEMBA1002777	401	R-HEMBA1002777	5933
	HEMBA1002779	F-HEMBA1002779	402	R-HEMBA1002779	5934
	HEMBA1002780	F-HEMBA1002780	403	R-HEMBA1002780	5935
25	HEMBA1002794	F-HEMBA1002794	404	R-HEMBA1002794	5936
	HEMBA1002801	F-HEMBA1002801	405	R-HEMBA1002801	5937
	HEMBA1002810	F-HEMBA1002810	406	R-HEMBA1002810	5938
	HEMBA1002816	F-HEMBA1002816	407	R-HEMBA1002816	5939
30	HEMBA1002818	F-HEMBA1002818	408	K ILLIDITIONEOIO	0000
	HEMBA1002826	F-HEMBA1002826	409	R-HEMBA1002826	5940
	HEMBA1002823	F-HEMBA1002833	410	R-HEMBA1002833	5941
	HEMBA1002850	F-HEMBA1002850	411	R-HEMBA1002850	5942
	HEMBA1002863	F-HEMBA1002863	412	R-HEMBA1002863	5943
35	HEMBA1002876	F-HEMBA1002876	413	R-HEMBA1002876	5944
	HEMBA1002886	F-HEMBA1002886	414	R-HEMBA1002886	5945
	HEMBA1002896	F-HEMBA1002896	415	R-HEMBA1002896	5946
	HEMBA1002991	F-HEMBA1002921	416	R-HEMBA1002921	5947
40	HEMBA1002924	F-HEMBA1002924	417	R-HEMBA1002924	5948
	HEMBA1002934		418	R-HEMBA1002934	5949
	HEMBA1002935	F-HEMBA1002935	419	R-HEMBA1002935	5950
	HEMBA1002937	F-HEMBA1002937	420	R-HEMBA1002937	5951
45	HEMBA1002939	F-HEMBA1002939	421	R-HEMBA1002939	5952
	HEMBA1002934	F-HEMBA1002944	422	R-HEMBA1002944	5953
	HEMBA1002941	F-HEMBA1002951	423	R-HEMBA1002951	5954
	HEMBA1002951	F-HEMBA1002954	423	R-HEMBA1002954	5955
50	HEMBA1002968	F-HEMBA1002968	424	R-HEMBA1002954	5956
50	HEMBA1002970	F-HEMBA1002970	425 426	R-HEMBA1002970	5957
	HEMBA1002970	F-HEMBA1002971	427	R-HEMBA1002971	5958
	HEMBA1002973	F-HEMBA1002971	427	R-HEMBA1002973	5959
	HEMBA1002973	F-HEMBA1002973	429	R-nnnnnnnnnnnn	5960
55	100mm1007991	1. IEMDV1007221	743	v innuminumin	5300

	HEMBA1002999	F-HEMBA1002999	430	R-HEMBA1002999	5961
	HEMBA1003021	F-HEMBA1003021	431	R-HEMBA1003021	5962
	HEMBA1003033	F-HEMBA1003033	432	R-HEMBA1003033	5963
5	HEMBA1003034	F-HEMBA1003034	433	R-HEMBA1003034	5964
	HEMBA1003035	F-HEMBA1003035	434	R-HEMBA1003035	5965
	HEMBA1003037	F-HEMBA1003037	435	R-HEMBA1003037	5966
	HEMBA1003041	F-HEMBA1003041	436	R-HEMBA1003041	5967
10	HEMBA1003046	F-HEMBA1003046	437	R-HEMBA1003046	5968
	HEMBA1003064	F-HEMBA1003064	438	R-HEMBA1003064	5969
	HEMBA1003067	F-HEMBA1003067	439	R-HEMBA1003067	5970
	HEMBA1003071	F-HEMBA1003071	440	R-HEMBA1003071	5971
15	HEMBA1003077	F-HEMBA1003077	441	R-HEMBA1003077	5972
	HEMBA1003078	F-HEMBA1003078	442	R-HEMBA1003078	5973
	HEMBA1003079	F-HEMBA1003079,	443	R-HEMBA1003079	5974
	HEMBA1003083	F-HEMBA1003083	444	R-HEMBA1003083	5975
20	HEMBA1003086	F-HEMBA1003086	445	R-HEMBA1003086	5976
20	HEMBA1003096	F-HEMBA1003096	446	R-HEMBA1003096	5977
	HEMBA1003098	F-HEMBA1003098	447	R-HEMBA1003098	5978
	HEMBA1003117	F-HEMBA1003117	448	R-HEMBA1003117	5979
	HEMBA1003129	F-HEMBA1003129	449	R-HEMBA1003129	5980
25	HEMBA1003133	F-HEMBA1003133	450	R-HEMBA1003133	5981
	HEMBA1003136	F-HEMBA1003136	451	R-HEMBA1003136	5982
	HEMBA1003142	F-HEMBA1003142	452	R-HEMBA1003142	5983
	HEMBA1003148	F-HEMBA1003148	453	R-HEMBA1003148	5984
30	HEMBA1003166	F-HEMBA1003166	454	R-HEMBA1003166	5985
	HEMBA1003175	F-HEMBA1003175	455	R-HEMBA1003175	5986
	HEMBA1003179	F-HEMBA1003179	456		
	HEMBA1003197	F-HEMBA1003197	457	R-HEMBA1003197	5987
35	HEMBA1003199	F-HEMBA1003199	458	R-HEMBA1003199	5988
	HEMBA1003202	F-HEMBA1003202	459	R-HEMBA1003202	5989
	HEMBA1003204	F-HEMBA1003204	460	R-HEMBA1003204	5990
	HEMBA1003212	F-HEMBA1003212	461	R-HEMBA1003212	5991
40	HEMBA1003220	F-HEMBA1003220	462	R-HEMBA1003220	5992
	HEMBA1003222	F-HEMBA1003222	463	R-HEMBA1003222	5993
	HEMBA1003229	F-HEMBA1003229	464	R-HEMBA1003229	5994
	HEMBA1003235	F-HEMBA1003235	465	R-HEMBA1003235	5995
	HEMBA1003250	F-HEMBA1003250	466	R-HEMBA1003250	5996
45	HEMBA1003257	F-HEMBA1003257	467	R-HEMBA1003257	5997
	HEMBA1003273		468	R-HEMBA1003273	5998
	HEMBA1003276		469	R-HEMBA1003276	5999
	HEMBA1003278		470	R-HEMBA1003278	6000
50	HEMBA1003281		471	R-HEMBA1003281	6001
	HEMBA1003286		472		
	HEMBA1003291		473	R-HEMBA1003291	6002
	HEMBA1003296		474	R-HEMBA1003296	6003
55	HEMBA1003304	F-HEMBA1003304	475	R-HEMBA1003304	6004

	HEMBA1003309	F-HEMBA1003309	476	R-HEMBA1003309	6005
	HEMBA1003314	F-HEMBA1003314	477	R-HEMBA1003314	6006
	HEMBA1003322	F-HEMBA1003322	478	R-HEMBA1003322	6007
5	HEMBA1003327	F-HEMBA1003327	479	R-HEMBA1003327	6008
	HEMBA1003328	F-HEMBA1003328	480	R-HEMBA1003328	6009
	HEMBA1003330	F-HEMBA1003330	481	R-HEMBA1003330	6010
	HEMBA1003348	F-HEMBA1003348	482	R-HEMBA1003348	6011
10	HEMBA1003369	F-HEMBA1003369	483	R-HEMBA1003369	6012
	HEMBA1003370	F-HEMBA1003370	484	R-HEMBA1003370	6013
	HEMBA1003373	F-HEMBA1003373	485	R-HEMBA1003373	6014
	HEMBA1003376	F-HEMBA1003376	486	R-HEMBA1003376	6015
15	HEMBA1003380	F-HEMBA1003380	487	R-HEMBA1003380	6016
15	HEMBA1003384	F-HEMBA1003384	488	R-HEMBA1003384	6017
	HEMBA1003395	F-HEMBA1003395,	489	R-HEMBA1003395	6018
	HEMBA1003402	F-HEMBA1003402	490	R-HEMBA1003402	6019
	HEMBA1003403	F-HEMBA1003403	491		
20	HEMBA1003408	F-HEMBA1003408	492	R-nnnnnnnnnnn	6020
	HEMBA1003417	F-HEMBA1003417	493	R-HEMBA1003417	6021
	HEMBA1003418	F-HEMBA1003418	494	R-HEMBA1003418	6022
	HEMBA1003433	F-HEMBA1003433	495	R-HEMBA1003433	6023
25	HEMBA1003447	F-HEMBA1003447	496		
	HEMBA1003461	F-HEMBA1003461	497	R-HEMBA1003461	6024
	HEMBA1003463	F-HEMBA1003463	498	R-HEMBA1003463	6025
	HEMBA1003480	F-HEMBA1003480	499	R-HEMBA1003480	6026
30	HEMBA1003528	F-HEMBA1003528	500	R-HEMBA1003528	6027
	HEMBA1003531	F-HEMBA1003531	501	R-HEMBA1003531	6028
	HEMBA1003538	F-HEMBA1003538	502	R-HEMBA1003538	6029
	HEMBA1003545	F-HEMBA1003545	503	R-HEMBA1003545	6030
35	HEMBA1003548	F-HEMBA1003548	504	R-HEMBA1003548	6031
	HEMBA1003555	F-HEMBA1003555	505	R-HEMBA1003555	6032
	HEMBA1003556	F-HEMBA1003556	506	R-HEMBA1003556	6033
	HEMBA1003560	F-HEMBA1003560	507	R-HEMBA1003560	6034
40	HEMBA1003568	F-HEMBA1003568	508	R-HEMBA1003568	6035
10	HEMBA1003569	F-HEMBA1003569	509	R-HEMBA1003569	6036
	HEMBA1003571	F-HEMBA1003571	510	R-HEMBA1003571	6037
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15	HEMBA1003711	F-HEMBA1003711	534	R-HEMBA1003711	6061
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	HEMBA1003715	F-HEMBA1003715	536	R-HEMBA1003715	6063
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	HEMBA1003827	F-HEMBA1003827	552		
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	HEMBA1003866	F-HEMBA1003866	557	R-HEMBA1003866	6083
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	HEMBA1004097	F-HEMBA1004097	591	R-HEMBA1004097	6117
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	HEMBA1004132	F-HEMBA1004132	594	R-HEMBA1004132	6119
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	HEMBA1004241	F-HEMBA1004241	610	R-HEMBA1004241	6135
	HEMBA1004246	F-HEMBA1004246	611	R-HEMBA1004246	6136
	HEMBA1004248	F-HEMBA1004248	612	R-HEMBA1004248	6137
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	HEMBA1004274	F-HEMBA1004274	616		
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	HEMBA1004276	F-HEMBA1004276	618	R-HEMBA1004276	6142
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	HEMBA1004864	F-HEMBA1004864	704	R-HEMBA1004864	6225
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				R-HEMBA1004944	6237
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	HEMBA1005075	F-HEMBA1005075	737	R-HEMBA1005075	6258
	HEMBA1005079	F-HEMBA1005079	738	R-HEMBA1005079	6259
40	HEMBA1005083	F-HEMBA1005083	739	R-HEMBA1005083	6260
	HEMBA1005101	F-HEMBA1005101	740	R-HEMBA1005101	6261
	HEMBA1005113	F-HEMBA1005113	741	R-HEMBA1005113	6262
	HEMBA1005123	F-HEMBA1005123	742	R-HEMBA1005123	6263
45	HEMBA1005133	F-HEMBA1005133	743	R-HEMBA1005133	6264
	HEMBA1005149	F-HEMBA1005149	744	R-HEMBA1005149	6265
	HEMBA1005152	F-HEMBA1005152	745	R-HEMBA1005152	6266
	HEMBA1005159	F-HEMBA1005159	746	R-HEMBA1005159	6267
50	HEMBA1005185	F-HEMBA1005185	747	R-HEMBA1005185	6268
	HEMBA1005201	F-HEMBA1005201	748	R-HEMBA1005201	6269
	HEMBA1005202	F-HEMBA1005202	749	R-HEMBA1005202	6270
	HEMBA1005206	F-HEMBA1005206	750		
	HEMBA1005219	F-HEMBA1005219	751	R-HEMBA1005219	6271

	HEMBA1005223	F-HEMBA1005223	752	R-HEMBA1005223	6272
	HEMBA1005232	F-HEMBA1005232	753	R-HEMBA1005232	6273 .
	HEMBA1005241	F-HEMBA1005241	754	R-HEMBA1005241	6274
5	HEMBA1005244	F-HEMBA1005244	755	R-HEMBA1005244	6275
3	HEMBA1005251	F-HEMBA1005251	756	R-HEMBA1005251	6276
	HEMBA1005252	F-HEMBA1005252	757	R-HEMBA1005252	6277
	HEMBA1005274	F-HEMBA1005274	758	R-HEMBA1005274	6278
	HEMBA1005275	F-HEMBA1005275	759	R-HEMBA1005275	6279
10	HEMBA1005293	F-HEMBA1005293	760	R-HEMBA1005293	6280
	HEMBA1005296	F-HEMBA1005296	761	R-HEMBA1005296	6281
		F-HEMBA1005304	762	R-HEMBA1005304	6282
	HEMBA1005304		763	R-HEMBA1005311	6283
15	HEMBA1005311	F-HEMBA1005311			6284
	HEMBA1005314	F-HEMBA1005314	764	R-HEMBA1005314	6285
	HEMBA1005315	F-HEMBA1005315,	765	R-HEMBA1005315	
	HEMBA1005318	F-HEMBA1005318	766	R-HEMBA1005318	6286
20	HEMBA1005331	F-HEMBA1005331	767	R-HEMBA1005331	6287
20	HEMBA1005338	F-HEMBA1005338	768	n .mmm	6000
	HEMBA1005353	F-HEMBA1005353	769	R-HEMBA1005353	6288
	HEMBA1005359	F-HEMBA1005359	770	R-HEMBA1005359	6289
	HEMBA1005367	F-HEMBA1005367	771	R-HEMBA1005367	6290
25	HEMBA1005372	F-HEMBA1005372	772	R-HEMBA1005372	6291
	HEMBA1005374	F-HEMBA1005374	773	R-HEMBA1005374	6292
	HEMBA1005382	F-HEMBA1005382	774		
	HEMBA1005389	F-HEMBA1005389	775	R-HEMBA1005389	6293
30	HEMBA1005394	F-HEMBA1005394	776	R-HEMBA1005394	6294
30	HEMBA1005403	F-HEMBA1005403	777	R-HEMBA1005403	6295
	HEMBA1005408	F-HEMBA1005408	778	R-HEMBA1005408	6296
	HEMBA1005410	F-HEMBA1005410	779	R-HEMBA1005410	6297
	HEMBA1005411	F-HEMBA1005411	780	R-HEMBA1005411	6298
35	HEMBA1005423	F-HEMBA1005423	781	R-HEMBA1005423	6299
	HEMBA1005426	F-HEMBA1005426	782	R-HEMBA1005426	6300
	HEMBA1005443	F-HEMBA1005443	783	R-HEMBA1005443	6301
	HEMBA1005447	F-HEMBA1005447	784	R-HEMBA1005447	6302
40	HEMBA1005468	F-HEMBA1005468	785	R-HEMBA1005468	6303
	HEMBA1005469	F-HEMBA1005469	786	R-HEMBA1005469	6304
	HEMBA1005472	F-HEMBA1005472	787	R-HEMBA1005472	6305
	HEMBA1005474	F-HEMBA1005474	788		
	HEMBA1005475	F-HEMBA1005475	789	R-HEMBA1005475	6306
45	HEMBA1005497	F-HEMBA1005497	790	R-HEMBA1005497	6307
	HEMBA1005500	F-HEMBA1005500	791	R-HEMBA1005500	6308
	HEMBA1005506	F-HEMBA1005506	792	R-HEMBA1005506	6309
	HEMBA1005508	F-HEMBA1005508	793	R-HEMBA1005508	6310
50	HEMBA1005511	F-HEMBA1005511	794	R-HEMBA1005511	6311
	HEMBA1005513	F-HEMBA1005513	795		
	HEMBA1005517	F-HEMBA1005517	796	R-HEMBA1005517	6312
		F-HEMBA1005518	797	R-HEMBA1005518	6313
	HEMBA1005518	L-UEWDW1009910	131	H HEIDATOOOTO	0010

	HEMBA1005520	F-HEMBA1005520	798	R-HEMBA1005520	6314
	HEMBA1005526	F-HEMBA1005526	799	R-HEMBA1005526	6315
	HEMBA1005528	F-HEMBA1005528	800	R-HEMBA1005528	6316
5	HEMBA1005530	F-HEMBA1005530	801	R-HEMBA1005530	6317
	HEMBA1005548	F-HEMBA1005548	802	R-HEMBA1005548	6318
	HEMBA1005552	F-HEMBA1005552	803	R-HEMBA1005552	6319
	HEMBA1005558	F-HEMBA1005558	804	R-HEMBA1005558	6320
10	HEMBA1005568	F-HEMBA1005568	805	R-HEMBA1005568	6321
,,	HEMBA1005570	F-HEMBA1005570	806	R-HEMBA1005570	6322
	HEMBA1005576	F-HEMBA1005576	807	R-HEMBA1005576	6323
	HEMBA1005577	F-HEMBA1005577		R-HEMBA1005577	6324
	HEMBA1005581	F-HEMBA1005581	809	R-HEMBA1005581	6325
15	HEMBA1005582	F-HEMBA1005582	810	R-HEMBA1005582	6326
	HEMBA1005583	F-HEMBA1005583,		R-HEMBA1005583	6327
	HEMBA1005588	F-HEMBA1005588	812	R-HEMBA1005588	6328
	HEMBA1005593	F-HEMBA1005593	813	R-HEMBA1005593	6329
20	HEMBA1005595	F-HEMBA1005595	814	R-HEMBA1005595	6330
	HEMBA1005606	F-HEMBA1005606	815	R-HEMBA1005606	6331
	HEMBA1005609	F-HEMBA1005609	816	R-HEMBA1005609	6332
	HEMBA1005616	F-HEMBA1005616	817	R-HEMBA1005616	6333
25	HEMBA1005621	F-HEMBA1005621	818	R-HEMBA1005621	6334
	HEMBA1005627	F-HEMBA1005627	819	R-HEMBA1005627	6335
	HEMBA1005631	F-HEMBA1005631	820	R-HEMBA1005631	6336
	HEMBA1005632	F-HEMBA1005632	821	R-HEMBA1005632	6337
30	HEMBA1005634	F-HEMBA1005634	822	R-HEMBA1005634	6338
	HEMBA1005666	F-HEMBA1005666	823	R-HEMBA1005666	6339
	HEMBA1005670	F-HEMBA1005670	824	R-HEMBA1005670	6340
	HEMBA1005679	F-HEMBA1005679	825	R-HEMBA1005679	6341
	HEMBA1005680	F-HEMBA1005680	826	R-HEMBA1005680	6342
35	HEMBA1005685	F-HEMBA1005685	827	R-HEMBA1005685	6343
	HEMBA1005699	F-HEMBA1005699	828	R-HEMBA1005699	6344
	HEMBA1005705	F-HEMBA1005705	829	R-HEMBA1005705	6345
	HEMBA1005717	F-HEMBA1005717	830	R-HEMBA1005717	6346
40	HEMBA1005732	F-HEMBA1005732	831	R-HEMBA1005732	6347
	HEMBA1005737	F-HEMBA1005737	832	R-HEMBA1005737	6348
	HEMBA1005746	F-HEMBA1005746	833	R-nnnnnnnnnnn	6349
	HEMBA1005755	F-HEMBA1005755	834	R-HEMBA1005755	6350
45	HEMBA1005765	F-HEMBA1005765	835	R-HEMBA1005765	6351
	HEMBA1005780	F-HEMBA1005780	836	R-HEMBA1005780	6352
	HEMBA1005813	F-HEMBA1005813	837	R-HEMBA1005813	6353
	HEMBA1005815	F-HEMBA1005815	838	R-HEMBA1005815	6354
50	HEMBA1005822	F-HEMBA1005822	839	R-HEMBA1005822	6355
	HEMBA1005829	F-HEMBA1005829	840	R-HEMBA1005829	6356 6357
	HEMBA1005834		841	R-HEMBA1005834	6358
	HEMBA1005852	F-HEMBA1005852	842	R-HEMBA1005852	
	HEMBA1005853	F-HEMBA1005853	843	R-HEMBA1005853	6359

	HEMBA1005884	F-HEMBA1005884	844	R-HEMBA1005884	6360
	HEMBA1005891	F-HEMBA1005891	845	R-HEMBA1005891	6361
	HEMBA1005894	F-HEMBA1005894	846	R-HEMBA1005894	6362
5	HEMBA1005909	F-HEMBA1005909	847	R-HEMBA1005909	6363
	HEMBA1005911	F-HEMBA1005911	848	R-HEMBA1005911	6364
	HEMBA1005921	F-HEMBA1005921	849	R-HEMBA1005921	6365
	HEMBA1005931	F-HEMBA1005931	850	R-HEMBA1005931	6366
10	HEMBA1005934	F-HEMBA1005934	851	R-HEMBA1005934	6367
10	HEMBA1005962	F-HEMBA1005962	852	R-HEMBA1005962	6368
	HEMBA1005963	F-HEMBA1005963	853	R-HEMBA1005963	6369
	HEMBA1005990	F-HEMBA1005990	854	R-HEMBA1005990	6370
	HEMBA1005991	F-HEMBA1005991	855	R-HEMBA1005991	6371
15	HEMBA1005999	F-HEMBA1005999	856	R-HEMBA1005999	6372
	HEMBA1006002	F-HEMBA1006002,		R-HEMBA1006002	6373
	HEMBA1006005	F-HEMBA1006005	858	R-HEMBA1006005	6374
	HEMBA1006031	F-HEMBA1006031	859	R-nnnnnnnnnnn	6375
20	HEMBA1006035	F-HEMBA1006035	860	R-HEMBA1006035	6376
	HEMBA1006036	F-HEMBA1006036	861	R-HEMBA1006036	6377
	HEMBA1006042	F-HEMBA1006042	862	R-HEMBA1006042	6378
	HEMBA1006067	F-HEMBA1006067	863	R-nnnnnnnnnnn	6379
25	HEMBA1006081	F-HEMBA1006081	864	R-HEMBA1006081	6380
	HEMBA1006090	F-HEMBA1006090	865	R-HEMBA1006090	6381
	HEMBA1006091	F-HEMBA1006091	866	R-HEMBA1006091	6382
	HEMBA1006100	F-HEMBA1006100	867	R-HEMBA1006100	6383
30	HEMBA1006108	F-HEMBA1006108	868	R-HEMBA1006108	6384
30	HEMBA1006121	F-HEMBA1006121	869	R-HEMBA1006121	6385
	HEMBA1006124	F-HEMBA1006124	870	R-HEMBA1006124	6386
	HEMBA1006130	F-HEMBA1006130	871	R-HEMBA1006130	6387
	HEMBA1006138	F-HEMBA1006138	872	R-nnnnnnnnnnn	6388
35	HEMBA1006142	F-HEMBA1006142	873	R-HEMBA1006142	6389
	HEMBA1006155	F-HEMBA1006155	874	R-HEMBA1006155	6390
	HEMBA1006158	F-HEMBA1006158	875	R-HEMBA1006158	6391
	HEMBA1006173	F-HEMBA1006173	876	R-HEMBA1006173	6392
40	HEMBA1006182	F-HEMBA1006182	877	R-HEMBA1006182	6393
	HEMBA1006198	F-HEMBA1006198	878	R-HEMBA1006198	6394
	HEMBA1006235	F-HEMBA1006235	879	R-HEMBA1006235	6395
	HEMBA1006248	F-HEMBA1006248	880	R-HEMBA1006248	6396
45	HEMBA1006252	F-HEMBA1006252	881	R-HEMBA1006252	6397
	HEMBA1006253	F-HEMBA1006253	882	R-HEMBA1006253	6398
	HEMBA1006259	F-HEMBA1006259	883	R-HEMBA1006259	6399
	HEMBA1006268	F-HEMBA1006268	884	R-HEMBA1006268	6400
	HEMBA1006272	F-HEMBA1006272	885	R-HEMBA1006272	6401
50	HEMBA1006278	F-HEMBA1006278	886	R-nnnnnnnnnnn	6402
	HEMBA1006283	F-HEMBA1006283	887	R-HEMBA1006283	6403
	HEMBA1006284	F-HEMBA1006284	888	R-HEMBA1006284	6404
	HEMBA1006291	F-HEMBA1006291	889	R-HEMBA1006291	6405

	HEMBA1006293	F-HEMBA1006293	890	R-HEMBA1006293	· 6406
	HEMBA1006309	F-HEMBA1006309	891	R-HEMBA1006309	6407
	HEMBA1006310	F-HEMBA1006310	892	R-HEMBA1006310	6408
5	HEMBA1006328	F-HEMBA1006328	893	R-HEMBA1006328	6409
	HEMBA1006334	F-HEMBA1006334	894	R-HEMBA1006334	6410
	HEMBA1006344	F-HEMBA1006344	895	R-HEMBA1006344	6411
	HEMBA1006347	F-HEMBA1006347	896	R-HEMBA1006347	6412
10	HEMBA1006349	F-HEMBA1006349	897	R-HEMBA1006349	6413
70	HEMBA1006359	F-HEMBA1006359	898	R-HEMBA1006359	6414
	HEMBA1006364	F-HEMBA1006364	899	R-HEMBA1006364	6415
	HEMBA1006377	F-HEMBA1006377	900	R-HEMBA1006377	6416
	HEMBA1006380	F-HEMBA1006380	901	R-HEMBA1006380	6417
15	HEMBA1006381	F-HEMBA1006381	902	R-HEMBA1006381	6418
	HEMBA1006398	F-HEMBA1006398,	903	R-HEMBA1006398	6419
	HEMBA1006416	F-HEMBA1006416	904	R-HEMBA1006416	6420
	HEMBA1006419	F-HEMBA1006419	905	R-HEMBA1006419	6421
20	HEMBA1006421	F-HEMBA1006421	906	R-HEMBA1006421	6422
	HEMBA1006424	F-HEMBA1006424	907	R-HEMBA1006424	6423
	HEMBA1006426	F-HEMBA1006426	908	R-HEMBA1006426	6424
•	HEMBA1006438	F-HEMBA1006438	909	R-HEMBA1006438	6425
25	HEMBA1006445	F-HEMBA1006445	910	R-HEMBA1006445	6426
-	HEMBA1006446	F-HEMBA1006446	911	R-HEMBA1006446	6427
	HEMBA1006461	F-HEMBA1006461	912	R-HEMBA1006461	6428
	HEMBA1006467	F-HEMBA1006467	913	R-HEMBA1006467	6429
20	HEMBA1006471	F-HEMBA1006471	914	R-HEMBA1006471	6430
30	HEMBA1006474	F-HEMBA1006474	915	R-HEMBA1006474	6431
	HEMBA1006483	F-HEMBA1006483	916	R-HEMBA1006483	6432
	HEMBA1006485	F-HEMBA1006485	917	R-HEMBA1006485	6433
	HEMBA1006486	F-HEMBA1006486	918	R-HEMBA1006486	6434
35	HEMBA1006489	F-HEMBA1006489	919	R-HEMBA1006489	6435
	HEMBA1006492	F-HEMBA1006492	920	R-HEMBA1006492	6436
	HEMBA1006494	F-HEMBA1006494	921	R-HEMBA1006494	6437
	HEMBA1006497	F-HEMBA1006497	922	R-HEMBA1006497	6438
40	HEMBA1006502	F-HEMBA1006502	923	R-HEMBA1006502	6439
	HEMBA1006507	F-HEMBA1006507	924	R-HEMBA1006507	6440
	HEMBA1006521	F-HEMBA1006521	925	R-HEMBA1006521	6441
	HEMBA1006530	F-HEMBA1006530	926	R-HEMBA1006530	6442
45	HEMBA1006535	F-HEMBA1006535	927	R-HEMBA1006535	6443
43	HEMBA1006540	F-HEMBA1006540	928	R-HEMBA1006540	6444
	HEMBA1006546	F-HEMBA1006546	929	R-HEMBA1006546	6445
	HEMBA1006559	F-HEMBA1006559	930	R-HEMBA1006559	6446
	HEMBA1006562	F-HEMBA1006562	931	R-HEMBA1006562	6447
50	HEMBA1006566	F-HEMBA1006566	932	R-HEMBA1006566	6448
	HEMBA1006569	F-HEMBA1006569	933	R-HEMBA1006569	6449
	HEMBA1006579		934	R-HEMBA1006579	6450
	HEMBA1006583	F-HEMBA1006583	935	R-HEMBA1006583	6451

	HEMBA1006595	F-HEMBA1006595	936	R-HEMBA1006595	6452
	HEMBA1006597	F-HEMBA1006597	937	R-HEMBA1006597	6453
	HEMBA1006612	F-HEMBA1006612	938	R-HEMBA1006612	6454
5	HEMBA1006617	F-HEMBA1006617	939	R-nnnnnnnnnnn	6455
	HEMBA1006624	F-HEMBA1006624	940	R-HEMBA1006624	6456
	HEMBA1006631	F-HEMBA1006631	941	R-HEMBA1006631	6457
	HEMBA1006635	F-HEMBA1006635	942	R-HEMBA1006635	6458
	HEMBA1006639	F-HEMBA1006639	943	R-HEMBA1006639	6459
10	HEMBA1006643	F-HEMBA1006643	944	R-HEMBA1006643	6460
u	HEMBA1006648	F-HEMBA1006648	945	R-HEMBA1006648	6461
	HEMBA1006652	F-HEMBA1006652	946	R-HEMBA1006652	6462
	HEMBA1006653	F-HEMBA1006653	947	R-HEMBA1006653	6463
15	HEMBA1006659	F-HEMBA1006659	948		
	HEMBA1006665	F-HEMBA1006665,	949	R-HEMBA1006665	6464
	HEMBA1006674	F-HEMBA1006674	950	R-HEMBA1006674	6465
	HEMBA1006676	F-HEMBA1006676	951	R-HEMBA1006676	6466
20	HEMBA1006682	F-HEMBA1006682	952	R-HEMBA1006682	6467
	HEMBA1006695	F-HEMBA1006695	953	R-HEMBA1006695	6468
	HEMBA1006696	F-HEMBA1006696	954	R-HEMBA1006696	6469
	HEMBA1006708	F-HEMBA1006708	955	R-HEMBA1006708	6470
25	HEMBA1006709	F-HEMBA1006709	956	R-HEMBA1006709	6471
23	HEMBA1006717	F-HEMBA1006717	957	R-HEMBA1006717	6472
	HEMBA1006737	F-HEMBA1006737	958	R-HEMBA1006737	6473
	HEMBA1006744	F-HEMBA1006744	959	R-HEMBA1006744	6474
	HEMBA1006754	F-HEMBA1006754	960	R-HEMBA1006754	6475
30	HEMBA1006758	F-HEMBA1006758	961	R-HEMBA1006758	6476
	HEMBA1006767	F-HEMBA1006767	962	R-HEMBA1006767	6477
	HEMBA1006779	F-HEMBA1006779	963	R-HEMBA1006779	6478
	HEMBA1006780	F-HEMBA1006780	964	R-HEMBA1006780	6479
35	HEMBA1006789	F-HEMBA1006789	965	R-HEMBA1006789	6480
	HEMBA1006795	F-HEMBA1006795	966	R-HEMBA1006795	6481
	HEMBA1006796	F-HEMBA1006796	967	R-HEMBA1006796	6482
	HEMBA1006807	F-HEMBA1006807	968	R-HEMBA1006807	6483
40	HEMBA1006821	F-HEMBA1006821	969	R-HEMBA1006821	6484
	HEMBA1006824	F-HEMBA1006824	970	R-HEMBA1006824	6485
	HEMBA1006832	F-HEMBA1006832	971	R-HEMBA1006832	6486
	HEMBA1006849	F-HEMBA1006849	972	R-HEMBA1006849	6487
	HEMBA1006865	F-HEMBA1006865	973	R-HEMBA1006865	6488
45	HEMBA1006877	F-HEMBA1006877	974	R-nnnnnnnnnnn	6489
	HEMBA1006885	F-HEMBA1006885	975	R-HEMBA1006885	6490
	HEMBA1006900	F-HEMBA1006900	976	R-HEMBA1006900	6491
	HEMBA1006914	F-HEMBA1006914	977		
50	HEMBA1006921	F-HEMBA1006921	978	R-HEMBA1006921	6492
	HEMBA1006926	F-HEMBA1006926	979	R-HEMBA1006926	6493
	HEMBA1006929	F-HEMBA1006929	980	R-HEMBA1006929	6494
	HEMBA1006936	F-HEMBA1006936	981	R-HEMBA1006936	6495

•	HEMBA1006938	F-HEMBA1006938	982	R-HEMBA1006938	6496
	HEMBA1006941	F-HEMBA1006941	983	R-HEMBA1006941	6497
	HEMBA1006949	F-HEMBA1006949	984	R-HEMBA1006949	6498
5	HEMBA1006973	F-HEMBA1006973	985	R-HEMBA1006973	6499
	HEMBA1006976	F-HEMBA1006976	986	R-HEMBA1006976	6500
	HEMBA1006993	F-HEMBA1006993	987	R-HEMBA1006993	6501
	HEMBA1006996	F-HEMBA1006996	988	R-HEMBA1006996	6502
40	HEMBA1007002	F-HEMBA1007002	989	R-HEMBA1007002	6503
10	HEMBA1007017	F-HEMBA1007017	990	R-HEMBA1007017	6504
	HEMBA1007018	F-HEMBA1007018	991	R-HEMBA1007018	6505
	HEMBA1007045	F-HEMBA1007045	992	R-HEMBA1007045	6506
	HEMBA1007051	F-HEMBA1007051	993	R-HEMBA1007051	6507
15	HEMBA1007052	F-HEMBA1007052	994	R-HEMBA1007052	6508
	HEMBA1007062	F-HEMBA1007062,	995	R-HEMBA1007062	6509
	HEMBA1007066	F-HEMBA1007066	996	R-HEMBA1007066	6510
	HEMBA1007073	F-HEMBA1007073	997	R-HEMBA1007073	6511
20	HEMBA1007078	F-HEMBA1007078	998	R-HEMBA1007078	6512
	HEMBA1007080	F-HEMBA1007080	999		
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	HEMBA1007087	F-HEMBA1007087	1001	R-HEMBA1007087	6514
25	HEMBA1007112	F-HEMBA1007112	1002	R-HEMBA1007112	6515
23	HEMBA1007113	F-HEMBA1007113	1003	R-HEMBA1007113	6516
	HEMBA1007121	F-HEMBA1007121	1004		
	HEMBA1007129	F-HEMBA1007129	1005	R-HEMBA1007129	6517
	HEMBA1007147	F-HEMBA1007147	1006	R-HEMBA1007147	6518
30	HEMBA1007149	F-HEMBA1007149	1007	R-HEMBA1007149	6519
	HEMBA1007151	F-HEMBA1007151	1008	R-HEMBA1007151	6520
	HEMBA1007174	F-HEMBA1007174	1009	R-nananananan	6521
	HEMBA1007178	F-HEMBA1007178	1010	R-HEMBA1007178	6522
35	HEMBA1007194	F-HEMBA1007194	1011	R-HEMBA1007194	6523
	HEMBA1007203	F-HEMBA1007203	1012	R-HEMBA1007203	6524
	HEMBA1007206	F-HEMBA1007206	1013	R-HEMBA1007206	6525
	HEMBA1007224	F-HEMBA1007224	1014	R-HEMBA1007224	6526
40	HEMBA1007243	F-HEMBA1007243	1015	·	
	HEMBA1007251	F-HEMBA1007251	1016	R-HEMBA1007251	6527
	HEMBA1007256	F-HEMBA1007256	1017	R-HEMBA1007256	6528
	HEMBA1007267	F-HEMBA1007267	1018	R-HEMBA1007267	6529
	HEMBA1007273	F-HEMBA1007273	1019	R-HEMBA1007273	6530
45	HEMBA1007279	F-HEMBA1007279	1020	R-HEMBA1007279	6531
	HEMBA1007281	F-HEMBA1007281	1021	R-HEMBA1007281	6532
	HEMBA1007288	F-HEMBA1007288	1022		6533
	HEMBA1007300	F-HEMBA1007300	1023		6534
50	HEMBA1007301	F-HEMBA1007301	1024		6535
	HEMBA1007319	F-HEMBA1007319	1025		6536
	HEMBA1007320		1026		6537
	HEMBA1007322	F-HEMBA1007322	1027	R-HEMBA1007322	6538

			1000		2500
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	HEMBA1007341	F-HEMBA1007341	1029	R-HEMBA1007341	6540
	HEMBA1007342	F-HEMBA1007342	1030	R-HEMBA1007342	6541
5	HEMBA1007347	F-HEMBA1007347	1031	R-HEMBA1007347	6542
	HEMBB1000005	F-HEMBB1000005	1032	R-HEMBB1000005	6543
	HEMBB1000008	F-HEMBB1000008	1033	R-HEMBB1000008	6544
	HEMBB1000018	F-HEMBB1000018	1034	R-HEMBB1000018	6545
10	HEMBB1000024	F-HEMBB1000024	1035	R-HEMBB1000024	6546
	HEMBB1000025	F-HEMBB1000025	1036	R-HEMBB1000025	6547
	HEMBB1000030	F-HEMBB1000030	1037	R-HEMBB1000030	6548
	HEMBB1000036	F-HEMBB1000036	1038	R-HEMBB1000036	6549
	HEMBB1000037	F-HEMBB1000037	1039	R-HEMBB1000037	6550
15	<b>НЕМВВ1000039</b>	F-HEMBB1000039	1040	R-HEMBB1000039	6551
	HEMBB1000044	F-HEMBB1000044,	1041	R-HEMBB1000044	6552
	HEMBB1000048	F-HEMBB1000048	1042	R-HEMBB1000048	6553
	HEMBB1000050	F-HEMBB1000050	1043	R-HEMBB1000050	6554
20	HEMBB1000054	F-HEMBB1000054	1044	R-HEMBB1000054	6555
	HEMBB1000055	F-HEMBB1000055	1045	R-HEMBB1000055	6556
	HEMBB1000059	F-HEMBB1000059	1046	R-HEMBB1000059	6557
	HEMBB1000083	F-HEMBB1000083	1047	R-HEMBB1000083	6558
25	HEMBB1000089	F-HEMBB1000089	1048	R-HEMBB1000089	6559
25	HEMBB1000099	F-HEMBB1000099	1049	R-HEMBB1000099	6560
	HEMBB1000103	F-HEMBB1000103	1050	R-HEMBB1000103	6561
	HEMBB1000113	F-HEMBB1000113	1051	R-HEMBB1000113	6562
	HEMBB1000119	F-HEMBB1000119	1052	R-HEMBB1000119	6563
30	HEMBB1000136	F-HEMBB1000136	1053	R-HEMBB1000136	6564
	HEMBB1000141	F-HEMBB1000141	1054	R-HEMBB1000141	6565
	HEMBB1000144	F-HEMBB1000144	1055	R-HEMBB1000144	6566
	HEMBB1000173	F-HEMBB1000173	1056	R-HEMBB1000173	6567
35	HEMBB1000175	F-HEMBB1000175	1057	R-HEMBB1000175	6568
	HEMBB1000198	F-HEMBB1000198	1058	R-HEMBB1000198	6569
	HEMBB1000215	F-HEMBB1000215	1059	R-HEMBB1000215	6570
	HEMBB1000217	F-HEMBB1000217	1060	R-HEMBB1000217	6571
	HEMBB1000218	F-HEMBB1000218	1061	R-HEMBB1000218	6572
40	HEMBB1000226	F-HEMBB1000226	1062	R-HEMBB1000226	6573
	HEMBB1000240	F-HEMBB1000240	1063	R-HEMBB1000240	6574
	HEMBB1000244	F-HEMBB1000244	1064	R-HEMBB1000244	6575
	HEMBB1000250	F-HEMBB1000250	1065	R-HEMBB1000250	6576
45	HEMBB1000258	F-HEMBB1000258	1066	R-HEMBB1000258	6577
	HEMBB1000264		1067	R-HEMBB1000264	6578
	HEMBB1000266		1068	R-HEMBB1000266	6579
	HEMBB1000272	-	1069	R-HEMBB1000272	6580
50	HEMBB1000274		1070	R-HEMBB1000274	6581
	HEMBB1000214	-	1071	R-HEMBB1000284	6582
	HEMBB1000284		1072	R-HEMBB1000307	6583
	HEMBB1000312	=	1073	R-HEMBB1000312	6584
	100001000312	1 110000010		.,	

	HEMBB1000317	F-HEMBB1000317	1074	R-HEMBB1000317	6585
	HEMBB1000318	F-HEMBB1000318	1075	R-HEMBB1000318	6586
	HEMBB1000335	F-HEMBB1000335	1076	R-HEMBB1000335	6587
5	HEMBB1000336	F-HEMBB1000336	1077	R-HEMBB1000336	6588
	HEMBB1000337	F-HEMBB1000337	1078	R-HEMBB1000337	6589
	HEMBB1000338	F-HEMBB1000338	1079	R-HEMBB1000338	6590
	HEMBB1000339	F-HEMBB1000339	1080	R-HEMBB1000339	6591
:10	HEMBB1000341	F-HEMBB1000341	1081	R-HEMBB1000341	6592
10	HEMBB1000343	F-HEMBB1000343	1082	R-HEMBB1000343	6593
	HEMBB1000354	F-HEMBB1000354	1083	R-HEMBB1000354	6594
	HEMBB1000369	F-HEMBB1000369	1084	R-HEMBB1000369	6595
	HEMBB1000374	F-HEMBB1000374	1085	R-HEMBB1000374	6596
15	HEMBB1000376	F-HEMBB1000376	1086	R-HEMBB1000376	6597
	HEMBB1000391	F-HEMBB1000391,	1087	R-HEMBB1000391	6598
	HEMBB1000399	F-HEMBB1000399	1088	R-HEMBB1000399	6599
	HEMBB1000402	F-HEMBB1000402	1089	R-HEMBB1000402	6600
20	HEMBB1000404	F-HEMBB1000404	1090	R-HEMBB1000404	6601
	HEMBB1000420	F-HEMBB1000420	1091	R-HEMBB1000420	6602
	HEMBB1000434	F-HEMBB1000434	1092	R-HEMBB1000434	6603
	HEMBB1000438	F-HEMBB1000438	1093	R-HEMBB1000438	6604
25	HEMBB1000441	F-HEMBB1000441	1094	R-HEMBB1000441	6605
	HEMBB1000449	F-HEMBB1000449	1095	R-HEMBB1000449	6606
	HEMBB1000455	F-HEMBB1000455	1096	R-HEMBB1000455	6607
	HEMBB1000472	F-HEMBB1000472	1097	R-HEMBB1000472	6608
00	HEMBB1000480	F-HEMBB1000480	1098	R-HEMBB1000480	6609
30	HEMBB1000487	F-HEMBB1000487	1099	R-HEMBB1000487	6610
	HEMBB1000490	F-HEMBB1000490	1100	R-HEMBB1000490	6611
	HEMBB1000491	F-HEMBB1000491	1101	R-HEMBB1000491	6612
	HEMBB1000493	F-HEMBB1000493	1102	R-HEMBB1000493	6613
35	HEMBB1000510	F-HEMBB1000510	1103	R-HEMBB1000510	6614
	HEMBB1000518	F-HEMBB1000518	1104	R-HEMBB1000518	6615
	HEMBB1000523	F-HEMBB1000523	1105	R-HEMBB1000523	6616
	HEMBB1000530	F-HEMBB1000530	1106	R-HEMBB1000530	6617
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	HEMBB1000554	F-HEMBB1000554	1108	R-HEMBB1000554	6619
	HEMBB1000556	F-HEMBB1000556	1109	R-HEMBB1000556	6620
	HEMBB1000564	F-HEMBB1000564	1110	R-HEMBB1000564	6621
45	HEMBB1000573	F-HEMBB1000573	1111	R-HEMBB1000573	6622
	HEMBB1000575	F-HEMBB1000575	1112	R-HEMBB1000575	6623
	HEMBB1000586	F-HEMBB1000586	1113	R-HEMBB1000586	6624
	HEMBB1000589	F-HEMBB1000589	1114	R-HEMBB1000589	6625
50	HEMBB1000591	F-HEMBB1000591	1115	R-HEMBB1000591	6626
50	HEMBB1000592	F-HEMBB1000592	1116	R-HEMBB1000592	6627
	HEMBB1000593	F-HEMBB1000593	1117	n urumniaaacaa	0000
	HEMBB1000598	F-HEMBB1000598	1118	R-HEMBB1000598	6628
	HEMBB1000623	F-HEMBB1000623	1119	R-HEMBB1000623	6629

	HEMBB1000630	F-HEMBB1000630	1120	R-HEMBB1000630	6630
	HEMBB1000631	F-HEMBB1000631	1121	R-HEMBB1000631	6631
	HEMBB1000632	F-HEMBB1000632	1122	R-HEMBB1000632	6632
5	HEMBB1000637	F-HEMBB1900637	1123	R-HEMBB1000637	6633
	HEMBB1000638	F-HEMBB1000638	1124	R-HEMBB1000638	6634
	HEMBB1000643	F-HEMBB1000643	1125	R-HEMBB1000643	6635
	HEMBB1000649	F-HEMBB1000649	1126	R-HEMBB1000649	6636
10	HEMBB1000652	F-HEMBB1000652	1127	R-HEMBB1000652	6637
	HEMBB1000665	F-HEMBB1000665	1128	R-HEMBB1000665	6638
	HEMBB1000671	F-HEMBB1000671	1129	R-HEMBB1000671	6639
	HEMBB1000673	F-HEMBB1000673	1130	R-HEMBB1000673	6640
15	HEMBB1000684	F-HEMBB1000684	1131	R-HEMBB1000684	6641
	HEMBB1000693	F-HEMBB1000693	1132	R-nnnnnnnnnnn	6642
	HEMBB1000705	F-HEMBB1000705	1133	R-HEMBB1000705	6643
	HEMBB1000706	F-HEMBB1000706	1134	R-HEMBB1000706	6644
00	HEMBB1000709	F-HEMBB1000709	1135	R-HEMBB1000709	6645
20	HEMBB1000725	F-HEMBB1000725	1136	R-HEMBB1000725	6646
	HEMBB1000726	F-HEMBB1000726	1137	R-HEMBB1000726	6647
	HEMBB1000738	F-HEMBB1000738	1138	R-HEMBB1000738	6648
	HEMBB1000749	F-HEMBB1000749	1139	R-HEMBB1000749	6649
25	HEMBB1000763	F-HEMBB1000763	1140	R-HEMBB1000763	6650
	HEMBB1000770	F-HEMBB1000770	1141	R-HEMBB1000770	6651
	HEMBB1000774	F-HEMBB1000774	1142		
	HEMBB1000781	F-HEMBB1000781	1143	R-HEMBB1000781	6652
30	HEMBB1000789	F-HEMBB1000789	1144	R-HEMBB1000789	6653
	HEMBB1000790	F-HEMBB1000790	1145	R-HEMBB1000790	6654
	HEMBB1000794	F-HEMBB1000794	1146	R-HEMBB1000794	6655
	HEMBB1000807	F-HEMBB1000807	1147	R-HEMBB1000807	6656
35	HEMBB1000810	F-HEMBB1000810	1148	R-HEMBB1000810	6657
	HEMBB1000821	F-HEMBB1000821	1149	R-HEMBB1000821	6658
	HEMBB1000822	F-HEMBB1000822	1150	R-HEMBB1000822	6659
	HEMBB1000826	F-HEMBB1000826	1151	R-HEMBB1000826	6660
40	HEMBB1000827	F-HEMBB1000827	1152	R-HEMBB1000827	6661
	HEMBB1000831	F-HEMBB1000831	1153	R-HEMBB1000831	6662
	HEMBB1000835	F-HEMBB1000835	1154	R-HEMBB1000835	6663
	HEMBB1000840	F-HEMBB1000840	1155	R-HEMBB1000840	6664
45	HEMBB1000848	F-HEMBB1000848	1156	R-HEMBB1000848	6665
75	HEMBB1000852	F-HEMBB1000852	1157	R-HEMBB1000852	6666
	HEMBB1000870	F-HEMBB1000870	1158	R-HEMBB1000870	6667
	HEMBB1000876	F-HEMBB1000876	1159	R-HEMBB1000876	6668
	HEMBB1000883	F-HEMBB1000883	1160	R-HEMBB1000883	6669
50	HEMBB1000887	F-HEMBB1000887	1161	R-HEMBB1000887	6670
	HEMBB1000888	F-HEMBB1000888	1162	R-HEMBB1000888	6671
	HEMBB1000890	F-HEMBB1000890	1163	R-HEMBB1000890	6672
	HEMBB1000893	F-HEMBB1000893	1164	R-HEMBB1000893	6673
55	HEMBB1000908	F-HEMBB1000908	1165	R-HEMBB1000908	6674

	HEMBB1000910	F-HEMBB1000910	1166	R-HEMBB1000910	6675
	HEMBB1000913	F-HEMBB1000913	1167	R-HEMBB1000913	6676
	HEMBB1000915	F-HEMBB1000915	1168	R-HEMBB1000915	6677
5	HEMBB1000917	F-HEMBB1000917	1169	R-HEMBB1000917	6678
	HEMBB1000927	F-HEMBB1000927	1170	R-HEMBB1000927	6679
	HEMBB1000947	F-HEMBB1000947	1171	R-HEMBB1000947	6680
	HEMBB1000959	F-HEMBB1000959	1172	R-HEMBB1000959	6681
10	HEMBB1000973	F-HEMBB1000973	1173	R-HEMBB1000973	6682
	HEMBB1000975	F-HEMBB1000975	1174	R-HEMBB1000975	6683
	HEMBB1000981	F-HEMBB1000981	1175	R-HEMBB1000981	6684
	HEMBB1000985	F-HEMBB1000985	1176	R-HEMBB1000985	6685
15	HEMBB1000991	F-HEMBB1000991	1177	R-HEMBB1000991	6686
	HEMBB1000996	F-HEMBB1000996	1178	R-HEMBB1000996	6687
	HEMBB1001004	F-HEMBB1001004,	1179	R-HEMBB1001004	6688
	HEMBB1001008	F-HEMBB1001008	1180	R-HEMBB1001008	6689
20	HEMBB1001011	F~HEMBB1001011	1181	R-HEMBB1001011	6690
20	HEMBB1001014	F-HEMBB1001014	1182	R-HEMBB1001014	6691
	HEMBB1001020	F-HEMBB1001020	1183	R-HEMBB1001020	6692
	HEMBB1001024	F-HEMBB1001024	1184	R-HEMBB1001024	6693
	HEMBB1001037	F-HEMBB1001037	1185	R-HEMBB1001037	6694
25	HEMBB1001047	F-HEMBB1001047	1186	R-HEMBB1001047	6695
	HEMBB1001051	F-HEMBB1001051	1187	R-HEMBB1001051	6696
	HEMBB1001056	F-HEMBB1001056	1188	R-HEMBB1001056	6697
	HEMBB1001058	F-HEMBB1001058	1189	R-HEMBB1001058	6698
30	HEMBB1001060	F-HEMBB1001060	1190	R-HEMBB1001060	6699
	HEMBB1001063	F-HEMBB1001063	1191	R-HEMBB1001063	6700
	HEMBB1001068	F-HEMBB1001068	1192	R-HEMBB1001068	6701
	HEMBB1001096	F-HEMBB1001096	1193	R-HEMBB1001096	6702
35	HEMBB1001102	F-HEMBB1001102	1194	R-HEMBB1001102	6703
	HEMBB1001105	F-HEMBB1001105	1195	R-HEMBB1001105	6704
	HEMBB1001112	F-HEMBB1001112	1196		
	HEMBB1001114	F-HEMBB1001114	1197	R-HEMBB1001114	6705
40	HEMBB1001117	F-HEMBB1001117	1198	R-HEMBB1001117	6706
	HEMBB1001119	F-HEMBB1001119	1199	R-HEMBB1001119	6707
	HEMBB1001126	F-HEMBB1001126	1200	R-HEMBB1001126	6708
	HEMBB1001133	F-HEMBB1001133	1201	R-HEMBB1001133	6709
45	HEMBB1001137	F-HEMBB1001137	1202	R-HEMBB1001137	6710
	HEMBB1001142	F-HEMBB1001142	1203	R-HEMBB1001142	6711
	HEMBB1001151	F-HEMBB1001151	1204	R-HEMBB1001151	6712
	HEMBB1001153	F-HEMBB1001153	1205	R-HEMBB1001153	6713
50	HEMBB1001169	F-HEMBB1001169	1206	R-HEMBB1001169	6714
	HEMBB1001175	F-HEMBB1001175	1207	R-nnnnnnnnnnn	6715
	HEMBB1001177	F-HEMBB1001177	1208	R-HEMBB1001177	6716
	HEMBB1001182	F-HEMBB1001182	1209	R-HEMBB1001182	6717
	HEMBB1001199	F-HEMBB1001199	1210	R-HEMBB1001199	6718
55	HEMBB1001208	F-HEMBB1001208	1211	R-HEMBB1001208	6719

	HEMBB1001209	F-HEMBB1001209	1212	R-HEMBB1001209	6720
	HEMBB1001210	F-HEMBB1001210	1213	R-HEMBB1001210	6721
	HEMBB1001218	F-HEMBB1001218	1214	R-HEMBB1001218	6722
5	HEMBB1001221	F-HEMBB1001221	1215	R-HEMBB1001221	6723
	HEMBB1001234	F-HEMBB1001234	1216	R-HEMBB1001234	6724
	HEMBB1001242	F-HEMBB1001242	1217	R-HEMBB1001242	6725
	HEMBB1001249	F-HEMBB1001249	1218	R-HEMBB1001249	6726
10	HEMBB1001253	F-HEMBB1001253	1219	R-HEMBB1001253	6727
	HEMBB1001254	F-HEMBB1001254	1220	R-HEMBB1001254	6728
	HEMBB1001267	F-HEMBB1001267	1221	R-HEMBB1001267	6729
	HEMBB1001271	F-HEMBB1001271	1222	R-HEMBB1001271	6730
15	HEMBB1001282	F-HEMBB1001282	1223	R-HEMBB1001282	6731
13	HEMBB1001288	F-HEMBB1001288	1224	R-HEMBB1001288	6732
	HEMBB1001289	F-HEMBB1001289,	1225	R-HEMBB1001289	6733
	HEMBB1001294	F-HEMBB1001294	1226	R-HEMBB1001294	6734
	HEMBB1001302	F-HEMBB1001302	1227	R-HEMBB1001302	6735
20	HEMBB1001304	F-HEMBB1001304	1228	R-HEMBB1001304	6736
	HEMBB1001314	F-HEMBB1001314	1229	R-HEMBB1001314	6737
	HEMBB1001315	F-HEMBB1001315	1230	R-HEMBB1001315	6738
	HEMBB1001317	F-HEMBB1001317	1231	R-HEMBB1001317	6739
25	HEMBB1001326	F-HEMBB1001326	1232	R-HEMBB1001326	6740
	HEMBB1001331	F-HEMBB1001331	1233	R-HEMBB1001331	6741
	HEMBB1001335	F-HEMBB1001335	1234	R-HEMBB1001335	6742
	HEMBB1001337	F-HEMBB1001337	1235	R-HEMBB1001337	6743
30	HEMBB1001339	F-HEMBB1001339	1236	R-HEMBB1001339	6744
	HEMBB1001346	F-HEMBB1001346	1237	R-HEMBB1001346	6745
	HEMBB1001348	F-HEMBB1001348	1238	R-HEMBB1001348	6746
	HEMBB1001356	F-HEMBB1001356	1239	R-HEMBB1001356	6747
35	HEMBB1001364	F-HEMBB1001364	1240	R-HEMBB1001364	6748
	HEMBB1001366	F-HEMBB1001366	1241	R-HEMBB1001366	6749
	HEMBB1001367	F-HEMBB1001367	1242	R-HEMBB1001367	6750
	HEMBB1001369	F-HEMBB1001369	1243	R-HEMBB1001369	6751
40	HEMBB1001380	F-HEMBB1001380	1244	R-HEMBB1001380	6752
	HEMBB1001384	F-HEMBB1001384	1245	R-HEMBB1001384	6753
	HEMBB1001387	F-HEMBB1001387	1246	R-HEMBB1001387	6754
	HEMBB1001394	F-HEMBB1001394	1247	R-HEMBB1001394	6755
45	HEMBB1001410	F-HEMBB1001410	1248	R-HEMBB1001410	6756
	HEMBB1001424	F-HEMBB1001424	1249	R-HEMBB1001424	6757
	HEMBB1001426	F-HEMBB1001426	1250	R-HEMBB1001426	6758
	HEMBB1001429	F-HEMBB1001429	1251	R-HEMBB1001429	6759
50	HEMBB1001436	F-HEMBB1001436	1252	R-HEMBB1001436	6760
	HEMBB1001443	F-HEMBB1001443	1253	R-HEMBB1001443	6761
	HEMBB1001449	F-HEMBB1001449	1254	R-HEMBB1001449	6762
	HEMBB1001454	F-HEMBB1001454	1255	R-HEMBB1001454	6763
55	HEMBB1001458	F-HEMBB1001458	1256	R-HEMBB1001458	6764
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	HEMBB1001464	F-HEMBB1001464	1258	R-HEMBB1001464	6766
	HEMBB1001482	F-HEMBB1001482	1259	R-HEMBB1001482	6767
	HEMBB1001500	F-HEMBB1001500	1260	R-HEMBB1001500	6768
5	HEMBB1001521	F-HEMBB1001521	1261	R-HEMBB1001521	6769
	HEMBB1001527	F-HEMBB1001527	1262	R-HEMBB1001527	6770
	HEMBB1001531	F-HEMBB1001531	1263	R-HEMBB1001531	6771
	HEMBB1001535	F-HEMBB1001535	1264	R-HEMBB1001535	6772
10	HEMBB1001536	F-HEMBB1001536	1265	R-HEMBB1001536	6773
	HEMBB1001537	F-HEMBB1001537	1266	R-HEMBB1001537	6774
	HEMBB1001555	F-HEMBB1001555	1267	R-HEMBB1001555	6775
	HEMBB1001562	F-HEMBB1001562	1268	R-HEMBB1001562	6776
15	HEMBB1001564	F-HEMBB1001564	1269	R-HEMBB1001564	6777
.5	HEMBB1001565	F-HEMBB1001565	1270	R-HEMBB1001565	6778
	HEMBB1001585	F-HEMBB1001585	1271	R-HEMBB1001585	6779
	HEMBB1001586	F-HEMBB1001586	1272	R-HEMBB1001586	6780
	HEMBB1001588	F-HEMBB1001588	1273	R-HEMBB1001588	6781
20 -	HEMBB1001603	F-HEMBB1001603	1274	R-HEMBB1001603	6782
	HEMBB1001618	F-HEMBB1001618	1275	R-HEMBB1001618	6783
	HEMBB1001619	F-HEMBB1001619	1276	R-HEMBB1001619	6784
	HEMBB1001630	F-HEMBB1001630	1277	R-HEMBB1001630	6785
25	HEMBB1001635	F-HEMBB1001635	1278	R-HEMBB1001635	6786
	HEMBB1001637	F-HEMBB1001637	1279	R-HEMBB1001637	6787
	HEMBB1001641	F-HEMBB1001641	1280	R-HEMBB1001641	6788
	HEMBB1001653	F-HEMBB1001653	1281	R-HEMBB1001653	6789
30	HEMBB1001665	F-HEMBB1001665	1282	R-HEMBB1001665	6790
•	HEMBB1001668	F-HEMBB1001668	1283	R-HEMBB1001668	6791
	HEMBB1001673	F-HEMBB1001673	1284	R-HEMBB1001673	6792
	HEMBB1001684	F-HEMBB1001684	1285	R-HEMBB1001684	6793
35	HEMBB1001685	F-HEMBB1001685	1286	R-HEMBB1001685	6794
	HEMBB1001695	F-HEMBB1001695	1287	R-HEMBB1001695	6795
	HEMBB1001704	F-HEMBB1001704	1288	R-HEMBB1001704	6796
	HEMBB1001706	F-HEMBB1001706	1289	R-HEMBB1001706	6797
40	HEMBB1001707	F-HEMBB1001707	1290	R-HEMBB1001707	6798
	HEMBB1001717	F-HEMBB1001717	1291	R-HEMBB1001717	6799
	HEMBB1001735	F-HEMBB1001735	1292	R-HEMBB1001735	6800
	HEMBB1001736	F-HEMBB1001736	1293	R-HEMBB1001736	6801
45	HEMBB1001747	F-HEMBB1001747	1294	R-HEMBB1001747	6802
	HEMBB1001749	F-HEMBB1001749	1295	R-HEMBB1001749	6803
	HEMBB1001753		1296	R-HEMBB1001753	6804
	HEMBB1001756		1297	R-HEMBB1001756	6805
50	HEMBB1001760		1298	R-HEMBB1001760	6806
50	HEMBB1001762		1299	R-HEMBB1001762	6807
	HEMBB1001785		1300	R-HEMBB1001785	6808
	HEMBB1001797		1301	R-HEMBB1001797	6809
	HEMBB1001802		1302	R-HEMBB1001802	6810
55	HEMBB1001812	F-HEMBB1001812	1303	R-HEMBB1001812	6811

	HEMBB1001816	F-HEMBB1001816	1304	R-HEMBB1001816	6812
	HEMBB1001831	F-HEMBB1001831	1305	R-HEMBB1001831	6813
	HEMBB1001834	F-HEMBB1001834	1306		
5	HEMBB1001836	F-HEMBB1001836	1307	R-HEMBB1001836	6814
	HEMBB1001839	F-HEMBB1001839	1308	R-HEMBB1001839	6815
	HEMBB1001850	F-HEMBB1001850	1309	R-HEMBB1001850	6816
	HEMBB1001863	F-HEMBB1001863	1310	R-HEMBB1001863	6817
10	HEMBB1001867	F-HEMBB1001867	1311	R-HEMBB1001867	6818
	HEMBB1001868	F-HEMBB1001868	1312	R-HEMBB1001868	6819
	HEMBB1001869	F-HEMBB1001869	1313	R-HEMBB1001869	6820
	HEMBB1001872	F-HEMBB1001872	1314	R-HEMBB1001872	6821
15	HEMBB1001874	F-HEMBB1001874	1315	R-HEMBB1001874	6822
	HEMBB1001875	F-HEMBB1001875	1316	R-HEMBB1001875	6823
	HEMBB1001880	F-HEMBB1001880,	1317	R-HEMBB1001880	6824
	HEMBB1001899	F-HEMBB1001899	1318	R-HEMBB1001899	6825
	HEMBB1001905	F-HEMBB1001905	1319	R-HEMBB1001905	6826
20	HEMBB1001906	F-HEMBB1001906	1320	R-HEMBB1001906	6827
	HEMBB1001908	F-HEMBB1001908	1321	R-HEMBB1001908	6828
	HEMBB1001910	F-HEMBB1001910	1322	R-HEMBB1001910	6829
	HEMBB1001911	F-HEMBB1001911	1323	R-HEMBB1001911	6830
25	HEMBB1001915	F-HEMBB1001915	1324	R-HEMBB1001915	6831
	HEMBB1001921	F-HEMBB1001921	1325	R-HEMBB1001921	6832
	HEMBB1001922	F-HEMBB1001922	1326	R-HEMBB1001922	6833
	HEMBB1001925	F-HEMBB1001925	1327	R-HEMBB1001925	6834
30	HEMBB1001930	F-HEMBB1001930	1328	R-HEMBB1001930	6835
	HEMBB1001944	F-HEMBB1001944	1329	R-HEMBB1001944	6836
	HEMBB1001945	F-HEMBB1001945	1330	R-HEMBB1001945	6837
	HEMBB1001947	F-HEMBB1001947	1331	R-HEMBB1001947	6838
35	HEMBB1001950	F-HEMBB1001950	1332	R-HEMBB1001950	6839
	HEMBB1001952	F-HEMBB1001952	1333	R-HEMBB1001952	6840
	HEMBB1001953	F-HEMBB1001953	1334	R-HEMBB1001953	6841
	HEMBB1001957	F-HEMBB1001957	1335	R-HEMBB1001957	6842
40	HEMBB1001962	F-HEMBB1001962	1336	R-HEMBB1001962	6843
	HEMBB1001967	F-HEMBB1001967	1337	R-HEMBB1001967	6844
	HEMBB1001973	F-HEMBB1001973	1338	R-HEMBB1001973	6845
	HEMBB1001983	F-HEMBB1001983	1339	R-HEMBB1001983	6846
45	HEMBB1001988		1340	R-HEMBB1001988	6847
	HEMBB1001990		1341	R-HEMBB1001990	6848
	HEMBB1001996		1342	R-HEMBB1001996	6849
	HEMBB1001997	F-HEMBB1001997	1343	R-HEMBB1001997	6850
50	HEMBB1002002		1344	R-HEMBB1002002	6851
<b>J</b> 0	HEMBB1002005		1345	R-HEMBB1002005	6852
-	HEMBB1002009		1346	R-HEMBB1002009	6853
	HEMBB1002015		1347	R-HEMBB1002015	6854
55	HEMBB1002042		1348	R-HEMBB1002042	6855 6856
55	HEMBB1002043	F-HEMBB1002043	1349	R-HEMBB1002043	0650

	HEMBB1002044	F-HEMBB1002044	1350	R-HEMBB1002044	6857
•	HEMBB1002045	F-HEMBB1002045	1351	R-HEMBB1002045	6858
	HEMBB1002049	F-HEMBB1002049	1352	R-HEMBB1002049	6859
5	HEMBB1002050	F-HEMBB1002050	1353	R-HEMBB1002050	6860
	HEMBB1002068	F-HEMBB1002068	1354	R-HEMBB1002068	6861
	HEMBB1002069	F-HEMBB1002069	1355	R-HEMBB1002069	6862
	HEMBB1002092	F-HEMBB1002092	1356	R-HEMBB1002092	6863
10	HEMBB1002094	F-HEMBB1002094	1357	R-HEMBB1002094	6864
	HEMBB1002115	F-HEMBB1002115	1358	R-HEMBB1002115	6865
	HEMBB1002134	F-HEMBB1002134	1359		
	HEMBB1002139	F-HEMBB1002139	1360	R-HEMBB1002139	6866
15	HEMBB1002142	F-HEMBB1002142	1361	R-HEMBB1002142	6867
15	HEMBB1002152	F-HEMBB1002152	1362	R-HEMBB1002152	6868
	HEMBB1002189	F-HEMBB1002189,	1363	R-HEMBB1002189	6869
	HEMBB1002190	F-HEMBB1002190	1364	R-HEMBB1002190	6870
	HEMBB1002193	F-HEMBB1002193	1365	R-HEMBB1002193	6871
20	HEMBB1002217	F-HEMBB1002217	1366	R-HEMBB1002217	6872
	HEMBB1002218	F-HEMBB1002218	1367	R-HEMBB1002218	6873
	HEMBB1002232	F-HEMBB1002232	1368	R-HEMBB1002232	6874
	HEMBB1002247	F-HEMBB1002247	1369	R-HEMBB1002247	6875
25	HEMBB1002249	F-HEMBB1002249	1370	R-HEMBB1002249	6876
	HEMBB1002254	F-HEMBB1002254	1371	R-HEMBB1002254	6877
	HEMBB1002255	F-HEMBB1002255	1372	R-HEMBB1002255	6878
	HEMBB1002266	F-HEMBB1002266	1373	R-HEMBB1002266	6879
30	HEMBB1002280	F-HEMBB1002280	1374	R-HEMBB1002280	6880
	HEMBB1002300	F-HEMBB1002300	1375	R-HEMBB1002300	6881
	HEMBB1002306	F-HEMBB1002306	1376	R-HEMBB1002306	6882
	HEMBB1002327	F-HEMBB1002327	1377	R-HEMBB1002327	6883
35	HEMBB1002329	F-HEMBB1002329	1378	R-HEMBB1002329	6884
	HEMBB1002340	F-HEMBB1002340	1379	R-HEMBB1002340	6885
	HEMBB1002342	F-HEMBB1002342	1380	R-HEMBB1002342	6886
	HEMBB1002358	F-HEMBB1002358	1381	R-HEMBB1002358	6887
40	HEMBB1002359	F-HEMBB1002359	1382	R-HEMBB1002359	6888
	HEMBB1002364	F-HEMBB1002364	1383	R-HEMBB1002364	6889
	HEMBB1002371	F-HEMBB1002371	1384	R-HEMBB1002371	6890
	HEMBB1002381	F-HEMBB1002381	1385	R-HEMBB1002381	6891
45	HEMBB1002383	F-HEMBB1002383	1386	R-HEMBB1002383	6892
45	HEMBB1002387	F-HEMBB1002387	1387	R-HEMBB1002387	6893
	HEMBB1002409	F-HEMBB1002409	1388		
	HEMBB1002415	F-HEMBB1002415	1389	R-HEMBB1002415	6894
	HEMBB1002425	F-HEMBB1002425	1390	R-HEMBB1002425	6895
50	HEMBB1002442		1391	R-HEMBB1002442	6896
	HEMBB1002453		1392	R-HEMBB1002453	6897
	HEMBB1002457		1393	R-HEMBB1002457	6898
	HEMBB1002458		1394	R-HEMBB1002458	6899
55	HEMBB1002477	F-HEMBB1002477	1395	R-HEMBB1002477	6900

	HEMBB1002489	F-HEMBB1002489	1396	R-HEMBB1002489	6901
	HEMBB1002492	F-HEMBB1002492	1397	R-HEMBB1002492	6902
	HEMBB1002495	F-HEMBB1002495	1398	R-HEMBB1002495	6903
5	HEMBB1002502	F-HEMBB1002502	1399	R-HEMBB1002502	6904
	HEMBB1002509	F-HEMBB1002509	1400	R-HEMBB1002509	6905
	HEMBB1002510	F-HEMBB1002510	1401	R-HEMBB1002510	6906
	HEMBB1002520	F-HEMBB1002520	1402	R-HEMBB1002520	6907
10	HEMBB1002522	F-HEMBB1002522	1403	R-HEMBB1002522	6908
	HEMBB1002531	F-HEMBB1002531	1404	R-HEMBB1002531	6909
	HEMBB1002534	F-HEMBB1002534	1405	R-HEMBB1002534	6910
	HEMBB1002545	F-HEMBB1002545	1406	R-HEMBB1002545	6911
15	HEMBB1002550	F-HEMBB1002550	1407	R-HEMBB1002550	6912
,,	HEMBB1002556	F-HEMBB1002556	1408	R-HEMBB1002556	6913
	HEMBB1002579	F-HEMBB1002579	1409	R-HEMBB1002579	6914
	HEMBB1002582	F-HEMBB1002582	1410	R-HEMBB1002582	6915
20	HEMBB1002590	F-HEMBB1002590	1411	R-HEMBB1002590	6916
20	HEMBB1002596	F-HEMBB1002596	1412	R-HEMBB1002596	6917
	HEMBB1002600	F-HEMBB1002600	1413	R-HEMBB1002600	6918
	HEMBB1002601	F-HEMBB1002601	1414	R-HEMBB1002601	6919
	HEMBB1002603	F-HEMBB1002603	1415	R-HEMBB1002603	6920
25	HEMBB1002607	F-HEMBB1002607	1416	R-HEMBB1002607	6921
	HEMBB1002610	F-HEMBB1002610	1417	R-HEMBB1002610	6922
	HEMBB1002613	F-HEMBB1002613	1418	R-HEMBB1002613	6923
	HEMBB1002614	F-HEMBB1002614	1419	R-HEMBB1002614	6924
30	HEMBB1002617	F-HEMBB1002617	1420	R-HEMBB1002617	6925
	HEMBB1002623	F-HEMBB1002623	1421	R-HEMBB1002623	6926
	HEMBB1002635	F-HEMBB1002635	1422	R-HEMBB1002635	6927
	HEMBB1002664	F-HEMBB1002664	1423	R-HEMBB1002664	6928
35	HEMBB1002677	F-HEMBB1002677	1424	R-HEMBB1002677	6929
	HEMBB1002683	F-HEMBB1002683	1425	R-HEMBB1002683	6930
	HEMBB1002684	F-HEMBB1002684	1426	R-HEMBB1002684	6931
	HEMBB1002686	F-HEMBB1002686	1427	R-HEMBB1002686	6932
40	HEMBB1002692	F-HEMBB1002692	1428	R-HEMBB1002692	6933
	HEMBB1002697	F-HEMBB1002697	1429	R-HEMBB1002697	6934
	HEMBB1002699	F-HEMBB1002699	1430	R-HEMBB1002699	6935
	HEMBB1002702	F-HEMBB1002702	1431	R-HEMBB1002702	6936
45	HEMBB1002705	F-HEMBB1002705	1432	R-HEMBB1002705	6937
	HEMBB1002712	F-HEMBB1002712	1433	R-HEMBB1002712	6938
	MAMMA1000009	F-MAMMA1000009	1434	R-MAMMA1000009	6939
	MAMMA1000019	F-MAMMA1000019	1435	R-MAMMA1000019	6940
50	MAMMA1000020	F-MAMMA1000020	1436	R-MAMMA1000020	6941
<b>J</b>	MAMMA1000025	F-MAMMA1000025	1437	R-MAMMA1000025	6942
	MAMMA1000043	F-MAMMA1000043	1438	R-MAMMA1000043	6943
	MAMMA1000045	F-MAMMA1000045	1439	R-MAMMA1000045	6944
	MAMMA1000055	F-MAMMA1000055	1440	R-MAMMA1000055	6945
55	MAMMA1000057	F-MAMMA1000057	1 <del>44</del> 1	R-MAMMA1000057	6946

	MAMMA1000069	F-MAMMA1000069	1442	R-MAMMA1000069	6947
	MAMMA1000084	F-MAMMA1000084	1443	R-MAMMA1000084	6948
	MAMMA1000085	F-MAMMA1000085	1444	R-MAMMA1000085	6949
5	MAMMA1000092	F-MAMMA1000092	1445	R-MAMMA1000092	6950
	MAMMA1000103	F-MAMMA1000103	1446	R-MAMMA1000103	6951
	MAMMA1000117	F-MAMMA1000117	1447	R-MAMMA1000117	6952
	MAMMA1000129	F-MAMMA1000129	1448	R-MAMMA1000129	6953
10	MAMMA1000133	F-MAMMA1000133	1449	R-MAMMA1000133	6954
	MAMMA1000134	F-MAMMA1000134	1450	R-MAMMA1000134	6955
	MAMMA1000139	F-MAMMA1000139	1451	R-MAMMA1000139	6956
	MAMMA1000143	F-MAMMA1000143	1452	R-MAMMA1000143	6957
15	MAMMA1000155	F-MAMMA1000155	1453	R-MAMMA1000155	6958
	MAMMA1000163	F-MAMMA1000163	1454	R-MAMMA1000163	6959
	MAMMA1000171	F-MAMMA1000171,	1455	R-MAMMA1000171	6960
	MAMMA1000173	F-MAMMA1000173	1456	R-MAMMA1000173	6961
	MAMMA1000175	F-MAMMA1000175	1457	R-MAMMA1000175	6962
20	MAMMA1000183	F-MAMMA1000183	1458	R-MAMMA1000183	6963
	MAMMA1000198	F-MAMMA1000198	1459	R-MAMMA1000198	6964
	MAMMA1000221	F-MAMMA1000221	1460	R-MAMMA1000221	6965
	MAMMA1000227	F-MAMMA1000227	1461	R-MAMMA1000227	6966
25	MAMMA1000241	F-MAMMA1000241	1462	R-MAMMA10Q0241	6967
	MAMMA1000251	F-MAMMA1000251	1463	R-MAMMA1000251	6968
	MAMMA1000254	F-MAMMA1000254	1464	R-MAMMA1000254	6969
	MAMMA1000257	F-MAMMA1000257	1465	R-MAMMA1000257	6970
30	MAMMA1000264	F-MAMMA1000264	1466	R-MAMMA1000264	6971
	MAMMA1000266	F-MAMMA1000266	1467	R-MAMMA1000266	6972
	MAMMA1000270	F-MAMMA1000270	1468	R-MAMMA1000270	6973
	MAMMA1000277	F-MAMMA1000277	1469	R-MAMMA1000277	6974
35	MAMMA1000278	F-MAMMA1000278	1470	R-MAMMA1000278	6975
	MAMMA1000279	F-MAMMA1000279	1471	R-MAMMA1000279	6976
1	MAMMA1000284	F-MAMMA1000284	1472	R-MAMMA1000284	6977
	MAMMA1000287	F-MAMMA1000287	1473	R-MAMMA1000287	6978
40	MAMMA1000302	F-MAMMA1000302	1474	R-MAMMA1000302	6979
	MAMMA1000307	F-MAMMA1000307	1475	R-MAMMA1000307	6980
	MAMMA1000309	F-MAMMA1000309	1476	R-MAMMA1000309	6981
	MAMMA1000312	F-MAMMA1000312	1477	R-MAMMA1000312	6982
45	MAMMA1000313	F-MAMMA1000313	1478	R-MAMMA1000313	6983
45	MAMMA1000331	F-MAMMA1000331	1479	R-MAMMA1000331	6984
	MAMMA1000339	F-MAMMA1000339	1480	R-MAMMA1000339	6985
	MAMMA1000340	F-MAMMA1000340	1481	R-MAMMA1000340	6986
	MAMMA1000348	F-MAMMA1000348	1482	R-MAMMA1000348	6987
50	MAMMA1000356		1483	R-MAMMA1000356	6988
	MAMMA1000360		1484	R-MAMMA1000360	6989
	MAMMA1000361	F-MAMMA1000361	1485	R-MAMMA1000361	6990
	MAMMA1000372		1486	R-MAMMA1000372	6991
55	MAMMA1000385	F-MAMMA1000385	1487	R-MAMMA1000385	6992

	- MAMMA1000388	F-MAMMA1000388	1488	R-MAMMA1000388	6993
	MAMMA1000395	F-MAMMA1000395	1489	R-MAMMA1000395	6994
	MAMMA1000402	F-MAMMA1000402	1490	R-MAMMA1000402	6995
5	MAMMA1000410	F-MAMMA1000410	1491	R-MAMMA1000410	6996
	MAMMA1000413	F-MAMMA1000413	1492	R-MAMMA1000413	6997
	MAMMA1000414	F-MAMMA1000414	1493	R-MAMMA1000414	6998
	MAMMA1000416	F-MAMMA1000416	1494	R-MAMMA1000416	6999
10	MAMMA1000421	F-MAMMA1000421	1495	R-MAMMA1000421	7000
	MAMMA1000422	F-MAMMA1000422	1496	R-MAMMA1000422	7001
	MAMMA1000423	F-MAMMA1000423	1497	R-MAMMA1000423	7002
	MAMMA1000424	F-MAMMA1000424	1498	R-MAMMA1000424	7003
15	MAMMA1000429	F-MAMMA1000429	1499	R-MAMMA1000429	7004
	MAMMA1000431	F-MAMMA1000431	1500	R-MAMMA1000431	7005
	MAMMA1000444	F-MAMMA1000444,	1501	R-MAMMA1000444	7006
	MAMMA1000446	F-MAMMA1000446	1502	R-MAMMA1000446	7007
20	MAMMA1000458	F-MAMMA1000458	1503	R-MAMMA1000458	7008
20	MAMMA1000468	F-MAMMA1000468	1504	R-MAMMA1000468	7009
	MAMMA1000472	F-MAMMA1000472	1505	R-MAMMA1000472	7010
	MAMMA1000478	F-MAMMA1000478	1506	R-MAMMA1000478	7011
	MAMMA1000483	F-MAMMA1000483	1507	R-MAMMA1000483	7012
25	MAMMA1000490	F-MAMMA1000490	1508	R-MAMMA1000490	7013
	MAMMA1000500	F-MAMMA1000500	1509	R-MAMMA1000500	7014
	MAMMA1000501	F-MAMMA1000501	1510	R-MAMMA1000501	7015
	MAMMA1000516	F-MAMMA1000516	1511	R-MAMMA1000516	7016
30	MAMMA1000522	F-MAMMA1000522	1512	R-MAMMA1000522	7017
	MAMMA1000524	F-MAMMA1000524	1513		
	MAMMA1000559	F-MAMMA1000559	1514	R-MAMMA1000559	7018
	MAMMA1000565	F-MAMMA1000565	1515	R-MAMMA1000565	7019
35	MAMMA1000567	F-MAMMA1000567	1516	R-MAMMA1000567	7020
	MAMMA1000576	F-MAMMA1000576	1517	R-MAMMA1000576	7021
	MAMMA1000583	F-MAMMA1000583	1518	R-MAMMA1000583	7022
	MAMMA1000585	F-MAMMA1000585	1519	R-MAMMA1000585	7023
40	MAMMA1000594	F-MAMMA1000594	1520	R-MAMMA1000594	7024
	MAMMA1000597	F-MAMMA1000597	1521	R-MAMMA1000597	7025
	MAMMA1000605	F-MAMMA1000605	1522	R-MAMMA1000605	7026
	MAMMA1000612	F-MAMMA1000612	1523	R-MAMMA1000612	7027
45	MAMMA1000616	F-MAMMA1000616	1524	R-MAMMA1000616	7028
	MAMMA1000621	F-MAMMA1000621	1525	R-MAMMA1000621	7029
	MAMMA1000623	F-MAMMA1000623	1526	R-MAMMA1000623	7030
	MAMMA1000625	F-MAMMA1000625	1527	R-MAMMA1000625	7031
50	MAMMA1000643	F-MAMMA1000643	1528	R-MAMMA1000643	7032
50	MAMMA1000664	F-MAMMA1000664	1529	R-MAMMA1000664	7033
	MAMMA1000669	F-MAMMA1000669	1530	R-MAMMA1000669	7034
	MAMMA1000670	F-MAMMA1000670	1531	R-MAMMA1000670	7035
	MAMMA1000672	F-MAMMA1000672	1532	R-MAMMA1000672	7036
55	MAMMA1000684	F-MAMMA1000684	1533	R-MAMMA1000684	7037

	MAMMA1000696	F-MAMMA1000696	1534	R-MAMMA1000696	7038
	MAMMA1000707	F-MAMMA1000707	1535	R-MAMMA1000707	7039
	MAMMA1000713	F-MAMMA1000713	1536	R-MAMMA1000713	7040
5	MAMMA1000714	F-MAMMA1000714	1537	R-MAMMA1000714	7041
	MAMMA1000718	F-MAMMA1000718	1538	R-MAMMA1000718	7042
	MAMMA1000720	F-MAMMA1000720	1539	R-MAMMA1000720	7043
	MAMMA1000723	F-MAMMA1000723	1540	R-MAMMA1000723	7044
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	MAMMA1000732	F-MAMMA1000732	1542	R-MAMMA1000732	7046
	MAMMA1000733	F-MAMMA1000733	1543	R-MAMMA1000733	7047
	MAMMA1000734	F-MAMMA1000734	1544	R-MAMMA1000734	7048
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	MAMMA1000744	F-MAMMA1000744	1546	R-MAMMA1000744	7050
	MAMMA1000746	F-MAMMA1000746,	1547	R-MAMMA1000746	7051
	MAMMA1000752	F-MAMMA1000752	1548	R-MAMMA1000752	7052
20	MAMMA1000760	F-MAMMA1000760	1549	R-MAMMA1000760	7053
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	MAMMA1000775	F-MAMMA1000775	1551	R-MAMMA1000775	7055
	MAMMA1000776	F-MAMMA1000776	1552	R-MAMMA1000776	7056
	MAMMA1000778	F-MAMMA1000778	1553	R-MAMMA1000778	7057
25	MAMMA1000782	F-MAMMA1000782	1554	R-MAMMA1000782	7058
	MAMMA1000798	F-MAMMA1000798	1555	R-MAMMA1000798	7059
	MAMMA1000802	F-MAMMA1000802	1556	R-MAMMA1000802	7060
	MAMMA 1000824	F-MAMMA1000824	1557		
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	MAMMA 1000839	F-MAMMA1000839	1559	R-MAMMA1000839	7062
	MAMMA 1000841	F-MAMMA1000841	1560	R-MAMMA1000841	7063
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	MAMMA1000845	F-MAMMA1000845	1563	R-MAMMA1000845	7066
	MAMMA1000851	F-MAMMA1000851	1564	R-MAMMA1000851	7067
	MAMMA1000855	F-MAMMA1000855	1565	R-MAMMA1000855	7068
40	MAMMA1000856	F-MAMMA1000856	1566	R-MAMMA1000856	7069
	MAMMA1000859	F-MAMMA1000859	1567	D WHIRM 10000CB	7070
	MAMMA1000862	F-MAMMA1000862	1568	R-MAMMA1000862 R-MAMMA1000863	7071
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	MAMMA1000867	F-MAMMA1000867	1571	R-MAMMA1000875	7074
-	MAMMA1000875	F-MAMMA1000875	1572 1573	R-MAMMA1000876	7075
	MAMMA1000876 MAMMA1000877	F-MAMMA1000876 F-MAMMA1000877	1574	R-MAMMA1000877	7076
50		F-MAMMA1000877	1574	R-MAMMA1000880	7077
	MAMMA1000880 MAMMA1000883		1576	R-MAMMA1000883	7078
	MAMMA 1000883	F-MAMMA1000883	1577	R-MAMMA1000897	7079
	MAMMA1000897 MAMMA1000905		1578	R-MAMMA1000905	7080
55	MAMMA 1000905 MAMMA 1000906	•	1579	R-MAMMA1000906	7081
55	MAMMA 1000300	L-MANATATION 300	1013	" unfilmention on o	. 501

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	MAMMA1000921	F-MAMMA1000921	1582	R-MAMMA1000921	7084
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	MAMMA1000940	F-MAMMA1000940	1584	R-MAMMA1000940	7086
	MAMMA1000941	F-MAMMA1000941	1585	R-MAMMA1000941	7087
	MAMMA1000942	F-MAMMA1000942	1586	R-MAMMA1000942	7088
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	MAMMA1000957	F-MAMMA1000957	1589	R-MAMMA1000957	7091
	MAMMA1000962	F-MAMMA1000962	1590	R-MAMMA1000962	7092
15	MAMMA1000968	F-MAMMA1000968	1591	R-MAMMA 1000968	7093
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	MAMMA1000979	F-MAMMA1000979,	1593	R-MAMMA1000979	7095
	MAMMA1000987	F-MAMMA1000987	1594	R-MAMMA1000987	7096
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	MAMMA1001021	F-MAMMA1001021	1598	R-MAMMA1001021	7100
	MAMMA1001024	F-MAMMA1001024	1599	R-MAMMA1001024	7101
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30	MAMMA1001050	F-MAMMA1001050	1604	R-MAMMA1001050	7106
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	MAMMA1001067	F-MAMMA1001067	1606	R-MAMMA1001067	7108
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35	MAMMA1001074	F-MAMMA1001074	1608	R-MAMMA1001074	7110
•	MAMMA1001075	F-MAMMA1001075	1609	R-MAMMA1001075	7111
	MAMMA1001078	F-MAMMA1001078	1610	R-MAMMA1001078	7112
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	MAMMA1001133	F-MAMMA1001133	1618	R-MAMMA1001133	7119
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	MAMMA1001162	F-MAMMA1001162	1624	R-MAMMA1001162	7125
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	MAMMA1001215	F-MAMMA1001215	1632	R-MAMMA1001215	7133
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	MAMMA1001243	F-MAMMA1001243	1635	R-MAMMA1001243	7136
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	MAMMA1001280	F-MAMMA1001280	1644	R-MAMMA1001280	7145
	MAMMA1001292	F-MAMMA1001292	1645	R-MAMMA1001292	7146
25	MAMMA1001296	F-MAMMA1001296	1646	R-MAMMA1001296	7147
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	MAMMA1001305	F-MAMMA1001305	1648	R-MAMMA1001305	7149
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30	MAMMA1001324	F-MAMMA1001324	1650	R-MAMMA1001324	7151
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	MAMMA1001341	F-MAMMA1001341	1652	R-MAMMA1001341	7153
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35	MAMMA1001346	F-MAMMA1001346	1654	R-MAMMA1001346	7155
	MAMMA1001383	F-MAMMA1001383	1655	R-MAMMA1001383	7156
	MAMMA1001388	F-MAMMA1001388	1656	R-MAMMA1001388	7157
	MAMMA1001397	F-MAMMA1001397	1657	R-MAMMA1001397	7158
40	MAMMA1001408	F-MAMMA1001408	1658	R-MAMMA1001408	7159
70	MAMMA1001411	F-MAMMA1001411	1659	R-MAMMA1001411	7160
	MAMMA1001419	F-MAMMA1001419	1660	R-MAMMA1001419	7161
	MAMMA1001420	F-MAMMA1001420	1661	R-MAMMA1001420	7162
	MAMMA1001435	F-MAMMA1001435	1662	R-MAMMA1001435	7163
45	MAMMA1001442	F-MAMMA1001442	1663	R-MAMMA1001442	7164
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	MAMMA1001452	F-MAMMA1001452	1665	R-MAMMA1001452	7166
	MAMMA1001465	F-MAMMA1001465	1666	R-MAMMA1001465	7167
50	MAMMA1001476	F-MAMMA1001476	1667	R-MAMMA1001476	7168
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	MAMMA1001501	F-MAMMA1001501	1669	R-MAMMA1001501	7170
	MAMMA1001502		1670	R-MAMMA1001502	7171
55	MAMMA1001510	F-MAMMA1001510	1671	R-MAMMA1001510	7172

	MAMMA1001522	F-MAMMA1001522	1672	R-MAMMA1001522	7173
	MAMMA1001547	F-MAMMA1001547	1673	R-MAMMA1001547	7174
	MAMMA1001551	F-MAMMA1001551	1674	R-MAMMA1001551	7175
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	MAMMA1001576	F-MAMMA1001576	1676	R-MAMMA1001576	7177
	MAMMA1001590	F-MAMMA1001590	1677	R-MAMMA1001590	7178
	MAMMA1001600	F-MAMMA1001600	1678	R-MAMMA1001600	7179
10	MAMMA1001604	F-MAMMA1001604	1679	R-MAMMA1001604	7180
	MAMMA1001606	F-MAMMA1001606	1680	R-MAMMA1001606	7181
	MAMMA1001620	F-MAMMA1001620	1681	R-MAMMA1001620	7182
	MAMMA1001627	F-MAMMA1001627	1682	R-MAMMA1001627	7183
15	MAMMA1001630	F-MAMMA1001630	1683	R-MAMMA1001630	7184
15	MAMMA1001633	F-MAMMA1001633	1684	R-MAMMA1001633	7185
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	MAMMA1001649	F-MAMMA1001649	1686	R-MAMMA1001649	7187
	MAMMA1001654	F-MAMMA1001654	1687		
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	MAMMA1001670	F-MAMMA1001670	1689	R-MAMMA1001670	7189
	MAMMA1001671	F-MAMMA1001671	1690	R-MAMMA1001671	7190
	MAMMA1001679	F-MAMMA1001679	1691	R-MAMMA1001679	7191
25	MAMMA1001683	F-MAMMA1001683	1692	R-MAMMA1001683	7192
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	MAMMA1001692	F-MAMMA1001692	1694	R-MAMMA1001692	7194
	MAMMA1001711	F-MAMMA1001711	1695	R-MAMMA1001711	7195
30	MAMMA1001715	F-MAMMA1001715	1696	R-MAMMA1001715	7196
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	MAMMA1001735	F-MAMMA1001735	1698	R-MAMMA1001735	7198
	MAMMA1001740	F-MAMMA1001740	1699	R-MAMMA1001740	7199
35	MAMMA1001743	F-MAMMA1001743	1700	R-MAMMA1001743	7200
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	MAMMA1001745	F-MAMMA1001745	1702	R-MAMMA1001745	7202
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40	MAMMA1001754	F-MAMMA1001754	1704	R-MAMMA1001754	7204
40	MAMMA1001757	F-MAMMA1001757	1705	R-MAMMA1001757	7205
	MAMMA1001760	F-MAMMA1001760	1706	R-MAMMA1001760	7206
	MAMMA1001764	F-MAMMA1001764	1707	R-MAMMA1001764	7207
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45	MAMMA1001769	F-MAMMA1001769	1709	R-MAMMA1001769	7209
	MAMMA1001771	F-MAMMA1001771	1710	R-MAMMA1001771	7210
	MAMMA1001783	F-MAMMA1001783	1711	R-MAMMA1001783	7211
	MAMMA1001785	F-MAMMA1001785	1712	R-MAMMA1001785	7212
50	MAMMA1001788	F-MAMMA1001788	1713	R-MAMMA1001788	7213
	MAMMA1001790	F-MAMMA1001790	1714	R-MAMMA1001790	7214
	MAMMA1001806	F-MAMMA1001806	1715	R-MAMMA1001806	7215
	MAMMA1001812	F-MAMMA1001812	1716	R-MAMMA1001812	7216
55	MAMMA1001815	F-MAMMA1001815	1717	R-MAMMA1001815	7217

	MAMMA1001817	F-MAMMA1001817	1718	R-MAMMA1001817	7218
	MAMMA1001818	F-MAMMA1001818	1719	R-MAMMA1001818	7219
	MAMMA1001820	F-MAMMA1001820	1720	R-MAMMA1001820	7220
5	MAMMA1001824	F-MAMMA1001824	1721	R-MAMMA1001824	7221
	MAMMA1001836	F-MAMMA1001836	1722	R-MAMMA1001836	7222
	MAMMA1001837	F-MAMMA1001837	1723	R-MAMMA1001837	7223
	MAMMA1001848	F-MAMMA1001848	1724	R-MAMMA1001848	7224
10	MAMMA1001851	F-MAMMA1001851	1725	R-MAMMA1001851	7225
	MAMMA1001854	F-MAMMA1001854	1726	R-MAMMA1001854	7226
	MAMMA1001858	F-MAMMA1001858	1727	R-MAMMA1001858	7227
	MAMMA1001864	F-MAMMA1001864	1728	R-MAMMA1001864	7228
15	MAMMA1001868	F-MAMMA1001868	1729	R-nnnnnnnnnnn	7229
73	MAMMA1001874	F-MAMMA1001874	1730	R-MAMMA1001874	7230
	MAMMA1001878	F-MAMMA1001878,	1731	R-MAMMA1001878	7231
	MAMMA1001880	F-MAMMA1001880	1732	R-MAMMA1001880	7232
	MAMMA1001890	F-MAMMA1001890	1733	R-MAMMA1001890	7233
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	MAMMA1001908	F-MAMMA1001908	1735	R-nnnnnnnnnnn	7235
	MAMMA1001931	F-MAMMA1001931	1736	R-MAMMA1001931	7236
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25	MAMMA1001963	F-MAMMA1001963	1738	R-MAMMA1001963	7238
	MAMMA1001969	F-MAMMA1001969	1739	R-MAMMA1001969	7239
	MAMMA1001970	F-MAMMA1001970	1740	R-MAMMA1001970	7240
	MAMMA1001992	F-MAMMA1001992	1741	R-MAMMA1001992	7241
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	MAMMA1002011	F-MAMMA1002011	1743	R-MAMMA1002011	7243
	MAMMA1002032	F-MAMMA1002032	1744	R-MAMMA1002032	7244
	MAMMA1002033	F-MAMMA1002033	1745	R-MAMMA1002033	7245
35	MAMMA1002041	F-MAMMA1002041	1746	R-MAMMA1002041	7246
	MAMMA1002042	F-MAMMA1002042	1747	R-MAMMA1002042	7247
	MAMMA1002047	F-MAMMA1002047	1748	R-MAMMA1002047	7248
	MAMMA1002056	F-MAMMA1002056	1749	R-MAMMA1002056	7249
40	MAMMA1002058	F-MAMMA1002058	1750	R-MAMMA1002058	7250
40	MAMMA1002068	F-MAMMA1002068	1751	R-MAMMA1002068	7251
	MAMMA1002078	F-MAMMA1002078	1752	R-MAMMA1002078	7252
	MAMMA1002082	F-MAMMA1002082	1753	R-MAMMA1002082	7253
	MAMMA1002084	F-MAMMA1002084	1754	R-MAMMA1002084	7254
45	MAMMA1002093	F-MAMMA1002093	1755	R-MAMMA1002093	7255
	MAMMA1002108	F-MAMMA1002108	1756	R-MAMMA1002108	7256
	MAMMA1002118	F-MAMMA1002118	1757	R-MAMMA1002118	7257
	MAMMA1002125	F-MAMMA1002125	1758	R-MAMMA1002125	7258
50	MAMMA1002132	F-MAMMA1002132	1759	R-MAMMA1002132	7259
	MAMMA1002140	F-MAMMA1002140	1760	R-MAMMA1002140	7260
	MAMMA1002143	F-MAMMA1002143	1761	R-MAMMA1002143	7261
	MAMMA1002145	F-MAMMA1002145	1762	R-MAMMA1002145	7262
55	MAMMA1002153	F-MAMMA1002153	1763	R-MAMMA1002153	7263

	MAMMA1002155	F-MAMMA1002155	1764	R-MAMMA1002155	7264
	MAMMA1002156	F-MAMMA1002156	1765	R-MAMMA1002156	7265
	MAMMA1002158	F-MAMMA1002158	1766	R-MAMMA1002158	7266
5	MAMMA1002170	F-MAMMA1002170	1767	R-MAMMA1002170	7267
	MAMMA1002174	F-MAMMA1002174	1768	R-MAMMA1002174	7268
	MAMMA1002198	F-MAMMA1002198	1769	R-MAMMA1002198	7269
	MAMMA1002209	F-MAMMA1002209	1770	R-MAMMA1002209	7270
10	MAMMA1002215	F-MAMMA1002215	1771	R-MAMMA1002215	7271
	MAMMA1002219	F-MAMMA1002219	1772	R-MAMMA1002219	7272
	MAMMA1002230	F-MAMMA1002230	1773	R-MAMMA1002230	7273
	MAMMA1002236	F-MAMMA1002236.	1774	R-MAMMA1002236	7274
15	MAMMA1002243	F-MAMMA1002243	1775	R-MAMMA1002243	7275
	MAMMA1002250	F-MAMMA1002250	1776	R-MAMMA1002250	7276
	MAMMA1002267	F-MAMMA1002267,	1777	R-MAMMA1002267	7277
	MAMMA1002268	F-MAMMA1002268	1778	R-MAMMA1002268	7278
20	MAMMA1002269	F-MAMMA1002269	1779	R-MAMMA1002269	7279
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	MAMMA1002299	F-MAMMA1002299	1786	R-MAMMA1002299	7286
	MAMMA1002308	F-MAMMA1002308	1787	R-MAMMA1002308	7287
30	MAMMA1002310	F-MAMMA1002310	1788	R-MAMMA1002310	7288
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	MAMMA1002312	F-MAMMA1002312	1790	R-MAMMA1002312	7290
	MAMMA1002317	F-MAMMA1002317	1791	R-MAMMA1002317	7291
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	MAMMA1002329	F-MAMMA1002329	1794	R-MAMMA1002329	7294
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15	MAMMA1002475	F-MAMMA1002475	1822	R-MAMMA1002475	7322
	MAMMA1002480	F-MAMMA1002480,	1823	R-MAMMA1002480	7323
	MAMMA1002485	F-MAMMA1002485	1824	R-MAMMA1002485	7324
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	MAMMA1002566	F-MAMMA1002566	1832	R-MAMMA1002566	7332
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,	MAMMA1002612	F-MAMMA1002612	1840	R-MAMMA1002612	7340
	MAMMA1002617	F-MAMMA1002617	1841	R-MAMMA1002617	7341
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	MAMMA1002622	F-MAMMA1002622	1844	R-MAMMA1002622	7344
	MAMMA1002623	F-MAMMA1002623	1845	R-MAMMA1002623	7345
	MAMMA1002625	F-MAMMA1002625	1846	R-MAMMA1002625	7346
45	MAMMA1002629	F-MAMMA1002629	1847	R-MAMMA1002629	7347
	MAMMA1002636	F-MAMMA1002636	1848	R-MAMMA1002636	7348
	MAMMA1002637	F-MAMMA1002637	1849	R-MAMMA1002637	7349
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	MAMMA1002662	F-MAMMA1002662	1853	R-MAMMA1002662 R-MAMMA1002665	7353 7354
	MAMMA1002665	F-MAMMA1002665	1854		
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	MAMMA1002685	F-MAMMA1002685	1858	R-MAMMA1002685	7358
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	MAMMA1002754	F-MAMMA1002754	1870	R-MAMMA1002754	7370
	MAMMA1002758	F-MAMMA1002758	1871	R-MAMMA1002758	7371
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	MAMMA1002769	F-MAMMA1002769	1874	R-MAMMA1002769	7374
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	MAMMA1002796	F-MAMMA1002796	1878	R-MAMMA1002796	7377
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45	MAMMA1002881	F-MAMMA1002881	1893	R-MAMMA1002881	7391
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55	MAMMA1002930	F-MAMMA1002930	1901	R-MAMMA1002930	7399

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_	MAMMA1002941	F-MAMMA1002941	1904	R-MAMMA1002941	7401
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	MAMMA1003026	F-MAMMA1003026	1919	R-MAMMA1003026	7415
	MAMMA1003031	F-MAMMA1003031	1920	R-MAMMA1003031	7416
25	MAMMA1003035	F-MAMMA1003035	1921	R-MAMMA1003035	7417
	MAMMA1003039	F-MAMMA1003039	1922	R-MAMMA1003039	7418
	MAMMA1003040	F-MAMMA1003040	1923	R-MAMMA1003040	7419
	MAMMA1003044	F-MAMMA1003044	1924	R-MAMMA1003044	7420
20	MAMMA1003047	F-MAMMA1003047	1925	R-MAMMA1003047	7421
30	MAMMA1003049	F-MAMMA1003049	1926	R-MAMMA1003049	7422
	MAMMA1003055	F-MAMMA1003055	1927	R-MAMMA1003055	7423
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	MAMMA1003127	F-MAMMA1003127	1935	R-MAMMA1003127	7431
	MAMMA1003135	F-MAMMA1003135	1936	R-MAMMA 1003135	7432
	MAMMA1003140	F-MAMMA1003140	1937	R-MAMMA1003140	7433
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	NT2RM1000035		1944		
	NT2RM1000037		1945		
55	NT2RM1000039		1946		
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	NT2RM2000030	F-NT2RM2000030	2045
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•	NT2RM2000092	F-NT2RM2000092	2048
	NT2RM2000093	F-NT2RM2000093	2049
	NT2RM2000101	F-NT2RM2000101	2050
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35	NT2RM2000407	F-NT2RM2000407	2068
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40	NT2RM2000469	F-NT2RM2000469	2072
	NT2RM2000490	F-NT2RM2000490	2073
	NT2RM2000502	F-NT2RM2000502	2074
	NT2RM2000504	F-NT2RM2000504	2075
45	NT2RM2000522	F-NT2RM2000522	2076 2077
,	NT2RM2000540 NT2RM2000556	F-NT2RM2000540 F-NT2RM2000556	2078
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	NT:	2RM2000837	F-NT2RM2000837	2103
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	NT	2RM2000952	F-NT2RM2000952	2105
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	NT	2RM2001004	F-NT2RM2001004	2107
	NT	2RM2001035	F-NT2RM2001035	2108
	NT	2RM2001065	F-NT2RM2001065	2109
	30 NT	2RM2001100	F-NT2RM2001100	2110
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		2RM2001196	F-NT2RM2001196	2117
		2RM2001201	F-NT2RM2001201	2118
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	-	T2RM2001256	F-NT2RM2001256	2123
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	- · ·	r2RM2001312	F-NT2RM2001312	2126
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15	NT2RM2001605	F-NT2RM2001605	2144
	NT2RM2001613	F-NT2RM2001613,	2145
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35	NT2RM2001695	F-NT2RM2001695	2160
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	NT2RM2001718	F-NT2RM2001718	2167
	NT2RM2001723	F-NT2RM2001723	2168
45	NT2RM2001727	F-NT2RM2001727	2169
	NT2RM2001730	F-NT2RM2001730	2170
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	NT2RM2001782		2176
55	NT2RM2001784	F-NT2RM2001784	2177

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30	NT2RM2002004	F-NT2RM2002004	2202		
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	NT2RM4000027	F-NT2RM4000027	2217	R-NT2RM4000027	7439
	NT2RM4000030	F-NT2RM4000030	2218	R-NT2RM4000030	7440
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	NT2RM4000085	F-NT2RM4000085	2221	R-NT2RM4000085	7443
	NT2RM4000086	F-NT2RM4000086	2222	R-NT2RM4000086	7444
	NT2RM4000104	F-NT2RM4000104	2223	R-NT2RM4000104	7445
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	NT2RM4000156	F-NT2RM4000156	2226	R-NT2RM4000156	7448
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	NT2RM4000191	F-NT2RM4000191	2229	R-NT2RM4000191	7451
	NT2RM4000197	F-NT2RM4000197	2230	R-NT2RM4000197	7452
	NT2RM4000199	F-NT2RM4000199	2231	R-NT2RM4000199	7453
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	NT2RM4000200 NT2RM4000202	F-NT2RM4000200		R-NT2RM4000200	7455
			2233	R-NT2RM4000202	7456
	NT2RM4000210	F-NT2RM4000210	2234		
15	NT2RM4000215	F-NT2RM4000215	2235	R-NT2RM4000215	7457
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	NT2RM4000290	F-NT2RM4000290	2241	R-NT2RM4000290	7463
	NT2RM4000324	F-NT2RM4000324	2242	R-NT2RM4000324	7464
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25	NT2RM4000344	F-NT2RM4000344	2244	R-NT2RM4000344	7466
	NT2RM4000349	F-NT2RM4000349	2245	R-NT2RM4000349	7467
	NT2RM4000354	F-NT2RM4000354	2246	R-NT2RM4000354	7468
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35	NT2RM4000421	F-NT2RM4000421	2253	R-NT2RM4000421	7475
	NT2RM4000425	F-NT2RM4000425	2254	R-NT2RM4000425	7476
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	NT2RM4000486	F-NT2RM4000486	2258	R-NT2RM4000486	7480
	NT2RM4000496	F-NT2RM4000496	2259	R-NT2RM4000496	7481
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.0	NT2RM4000515	F-NT2RM4000515	2262	R-nnnnnnnnnnn	7484
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	NT2RM4000531	F-NT2RM4000531	2264	R-NT2RM4000531	7486
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	NT2RM4000590	F-NT2RM4000590	2268	R-NT2RM4000590	7490
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	NT2RM4000616	F-NT2RM4000616	2272	R-NT2RM4000616	7494
5	NT2RM4000674	F-NT2RM4000674	2273	R-NT2RM4000674	7495
	NT2RM4000689	F-NT2RM4000689	2274	R-NT2RM4000689	7496
	NT2RM4000698	F-NT2RM4000698	2275	R-NT2RM4000698	7497
	NT2RM4000700	F-NT2RM4000700	2276	R-nnnnnnnnnnn	7498
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70	NT2RM4000717	F-NT2RM4000717	2278	R-NT2RM4000717	7500
	NT2RM4000733	F-NT2RM4000733	2279	R-NT2RM4000733	7501
	NT2RM4000734	F-NT2RM4000734	2280	R-NT2RM4000734	7502
	NT2RM4000741	F-NT2RM4000741	2281	R-NT2RM4000741	7503
15	NT2RM4000751	F-NT2RM4000751	2282	R-NT2RM4000751	7504
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25	NT2RM4000798	F-NT2RM4000798	2290	R-NT2RM4000798	7512
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	NT2RM4000833	F-NT2RM4000833	2293	R-NT2RM4000833	7515
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	NT2RM4000855	F-NT2RM4000855	2296	R-NT2RM4000855	7518
	NT2RM4000887	F-NT2RM4000887	2297	R-nnnnnnnnnnnn	7519
35	NT2RM4000895	F-NT2RM4000895	2298	R-NT2RM4000895	7520
55	NT2RM4000950	F-NT2RM4000950	2299	R-NT2RM4000950	7521
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40	NT2RM4001002	F-NT2RM4001002	2303	R-NT2RM4001002	7525
	NT2RM4001016	F-NT2RM4001016	2304	R-NT2RM4001016	7526
	NT2RM4001032	F-NT2RM4001032	2305	R-NT2RM4001032	7527
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45	NT2RM4001054	F-NT2RM4001054	2307	R-NT2RM4001054	7529
	NT2RM4001084	F-NT2RM4001084	2308	R-nnnnnnnnnn	7530
	NT2RM4001092	F-NT2RM4001092	2309	R-NT2RM4001092	7531
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	NT2RM4001155	F-NT2RM4001155	2313	R-NT2RM4001155	7535
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	NT2RM4001256	F-NT2RM4001256	2321	R-NT2RM4001256	7543
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	NT2RM4001340	F-NT2RM4001340	2327	R-NT2RM4001340	7549
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	NT2RM4001371	F-NT2RM4001371	2330	R-NT2RM4001371	7552
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	NT2RM4001412	F-NT2RM4001412	2335	R-NT2RM4001412	7557
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	NT2RM4001444	F-NT2RM4001444	2338	R-NT2RM4001444	7560
	NT2RM4001454	F-NT2RM4001454	2339	R-NT2RM4001454	7561
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30	NT2RM4001483	F-NT2RM4001483	2341	R-NT2RM4001483	7563
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	NT2RM4001519	F-NT2RM4001519	2343	R-NT2RM4001519	7565
	NT2RM4001522	F-NT2RM4001522	2344	R-NT2RM4001522	7566
35	NT2RM4001557	F-NT2RM4001557	2345	R-NT2RM4001557	7567
	NT2RM4001565	F-NT2RM4001565	2346	R-NT2RM4001565	7568
	NT2RM4001566	F-NT2RM4001566	2347	R-NT2RM4001566	7569
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	NT2RM4001592	F-NT2RM4001592	2350	R-nnnnnnnnnnn	7572
	NT2RM4001594	F-NT2RM4001594	2351	R-NT2RM4001594	7573
	NT2RM4001597	F-NT2RM4001597	2352	R-NT2RM4001597	7574
45	NT2RM4001605	F-NT2RM4001605	2353	R-NT2RM4001605	7575
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	NT2RM4001650	F-NT2RM4001650	2356	R-NT2RM4001650	7578
	NT2RM4001662	F-NT2RM4001662	2357	R-NT2RM4001662	7579
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	NT2RM4001710	F-NT2RM4001710	2360	R-NT2RM4001710	7582
	NT2RM4001714	F-NT2RM4001714	2361	R-NT2RM4001714	7583

	NT2RM4001715	F-NT2RM4001715	2362	R-nnnnnnnnnnn	7584
	NT2RM4001731	F-NT2RM4001731	2363	R-NT2RM4001731	7585
	NT2RM4001741	F-NT2RM4001741	2364	R-NT2RM4001741	7586
5	NT2RM4001746	F-NT2RM4001746	2365	R-NT2RM4001746	7587
	NT2RM4001754	F-NT2RM4001754	2366	R-NT2RM4001754	7588
	NT2RM4001758	F-NT2RM4001758	2367	R-NT2RM4001758	7589
	NT2RM4001776	F-NT2RM4001776	2368	R-NT2RM4001776	7590
10	NT2RM4001783	F-NT2RM4001783	2369	R-NT2RM4001783	7591
10	NT2RM4001810	F-NT2RM4001810	2370	R-NT2RM4001810	7592
	NT2RM4001813	F-NT2RM4001813	2371	R-NT2RM4001813	7593
	NT2RM4001819	F-NT2RM4001819	2372		
	NT2RM4001823	F-NT2RM4001823	2373	R-NT2RM4001823	7594
15	NT2RM4001828	F-NT2RM4001828	2374	R-NT2RM4001828	7595
	NT2RM4001836	F-NT2RM4001836,	2375	R-NT2RM4001836	7596
	NT2RM4001841	F-NT2RM4001841	2376	R-NT2RM4001841	7597
	NT2RM4001842	F-NT2RM4001842	2377	R-NT2RM4001842	7598
20	NT2RM4001856	F-NT2RM4001856	2378	R-NT2RM4001856	7599
	NT2RM4001858	F-NT2RM4001858	2379	R-nnnnnnnnnnn	7600
	NT2RM4001865	F-NT2RM4001865	2380	R-NT2RM4001865	7601
	NT2RM4001876	F-NT2RM4001876	2381	R-NT2RM4001876	7602
25	NT2RM4001880	F-NT2RM4001880	2382	R-NT2RM4001880	7603
20	NT2RM4001905	F-NT2RM4001905	2383	R-NT2RM4001905	7604
	NT2RM4001922	F-NT2RM4001922	2384	R-NT2RM4001922	7605
	NT2RM4001930	F-NT2RM4001930	2385	R-NT2RM4001930	7606
	NT2RM4001938	F-NT2RM4001938	2386	R-NT2RM4001938	7607
30	NT2RM4001940	F-NT2RM4001940	2387	R-NT2RM4001940	7608
	NT2RM4001953	F-NT2RM4001953	2388	R-NT2RM4001953	7609
	NT2RM4001965	F-NT2RM4001965	2389	R-NT2RM4001965	7610
	NT2RM4001969	F-NT2RM4001969	2390	R-nnnnnnnnnnn	7611
35	NT2RM4001979	F-NT2RM4001979	2391	R-NT2RM4001979	7612
	NT2RM4001984	F-NT2RM4001984	2392	R-NT2RM4001984	7613
	NT2RM4001987	F-NT2RM4001987	2393	R-NT2RM4001987	7614
	NT2RM4002013	F-NT2RM4002013	2394	R-NT2RM4002013	7615
40	NT2RM4002018	F-NT2RM4002018	2395	R-NT2RM4002018	7616
	NT2RM4002034	F-NT2RM4002034	2396	R-NT2RM4002034	7617
	NT2RM4002044	F-NT2RM4002044	2397	R-NT2RM4002044	7618
	NT2RM4002054	F-NT2RM4002054	2398	R-NT2RM4002054	7619
45	NT2RM4002055	F-NT2RM4002055	2399		
	NT2RM4002062		2400	R-NT2RM4002062	7620
	NT2RM4002063		2401	R-NT2RM4002063	7621
	NT2RM4002066		2402	R-nnnnnnnnnnn	7622
	NT2RM4002067		2403	R-NT2RM4002067	7623
50	NT2RM4002073		2404	R-NT2RM4002073	7624
	NT2RM4002075	••	2405	R-NT2RM4002075	7625
	NT2RM4002093		2406	R-NT2RM4002093	7626
	NT2RM4002109	F-NT2RM4002109	2407	R-nnnnnnnnnnn	7627

	NT2RM4002128	F-NT2RM4002128	2408	R-NT2RM4002128	7628
	NT2RM4002140	F-NT2RM4002140	2409	R-NT2RM4002140	7629
	NT2RM4002145	F-NT2RM4002145	2410	R-NT2RM4002145	7630
5	NT2RM4002146	F-NT2RM4002146	2411	R-NT2RM4002146	7631
	NT2RM4002161	F-NT2RM4002161	2412	R-NT2RM4002161	7632
	NT2RM4002174	F-NT2RM4002174	2413	R-NT2RM4002174	7633
	NT2RM4002189	F-NT2RM4002189	2414	R-NT2RM4002189	7634
10	NT2RM4002194	F-NT2RM4002194	2415	R-NT2RM4002194	7635
70	NT2RM4002205	F-NT2RM4002205	2416	R-NT2RM4002205	7636
	NT2RM4002213	F-NT2RM4002213	2417	R-NT2RM4002213	7637
	NT2RM4002226	F-NT2RM4002226	2418	R-NT2RM4002226	7638
	NT2RM4002251	F-NT2RM4002251	2419	R-NT2RM4002251	7639
15	NT2RM4002256	F-NT2RM4002256	2420	R-NT2RM4002256	7640
	NT2RM4002266	F-NT2RM4002266	2421	R-NT2RM4002266	7641
	NT2RM4002278	F-NT2RM4002278	2422	R-NT2RM4002278	7642
	NT2RM4002281	F-NT2RM4002281	2423	R-NT2RM4002281	7643
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	NT2RM4002294	F-NT2RM4002294	2425	R-NT2RM4002294	7645
	NT2RM4002301	F-NT2RM4002301	2426	R-NT2RM4002301	7646
	NT2RM4002323	F-NT2RM4002323	2427	R-NT2RM4002323	7647
25	NT2RM4002339	F-NT2RM4002339	2428	R-nnnnnnnnnnn	7648
	NT2RM4002344	F-NT2RM4002344	2429	R-NT2RM4002344	7649
	NT2RM4002373	F-NT2RM4002373	2430	R-NT2RM4002373	7650
	NT2RM4002374	F-NT2RM4002374	2431	R-NT2RM4002374	7651
	NT2RM4002383	F-NT2RM4002383	2432	R-NT2RM4002383	7652
30	NT2RM4002390	F-NT2RM4002390	2433	R-NT2RM4002390	7653
	NT2RM4002398	F-NT2RM4002398	2434		
	NT2RM4002409	F-NT2RM4002409	2435	R-NT2RM4002409	7654
	NT2RM4002438	F-NT2RM4002438	2436	R-NT2RM4002438	7655
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	NT2RM4002452	F-NT2RM4002452	2438	R-NT2RM4002452	7657
	NT2RM4002457	F-NT2RM4002457	2439	R-NT2RM4002457	7658
	NT2RM4002460	F-NT2RM4002460	2440	R-NT2RM4002460	7659
40	NT2RM4002479	F-NT2RM4002479	2441	R-NT2RM4002479	7660
•	NT2RM4002482	F-NT2RM4002482	2442	R-NT2RM4002482	7661
	NT2RM4002493	F-NT2RM4002493	2443	R-NT2RM4002493	7662
	NT2RM4002499	F-NT2RM4002499	2444	R-NT2RM4002499	7663
45	NT2RM4002504	F-NT2RM4002504	2445	R-NT2RM4002504	7664
40	NT2RM4002527	F-NT2RM4002527	2446	R-nnnnnnnnnnn	7665
	NT2RM4002532	F-NT2RM4002532	2447	R-NT2RM4002532	7666
	NT2RM4002534		2448	R-NT2RM4002534	7667
	NT2RM4002558		2449		
50	NT2RM4002565	F-NT2RM4002565	2450		
	NT2RM4002567		2451	R-NT2RM4002567	7668
	NT2RM4002571		2452	R-NT2RM4002571	7669
	NT2RM4002593	F-NT2RM4002593	2453	R-NT2RM4002593	7670

NT2RM4002623		NT2RM4002594	F-NT2RM4002594	2454		
NTZRP100035 F-NTZRP100035   2457     NTZRP1000060 F-NTZRP1000060   2458     NTZRP1000063 F-NTZRP1000060   2459     NTZRP10000101 F-NTZRP10000101   2461     NTZRP1000111 F-NTZRP1000111   2462     NTZRP1000112 F-NTZRP1000112   2463     NTZRP1000124 F-NTZRP1000124   2464     NTZRP1000130 F-NTZRP1000130   2465     NTZRP1000170 F-NTZRP1000170   2467     NTZRP1000170 F-NTZRP1000170   2467     NTZRP1000171 F-NTZRP1000171   2468     NTZRP1000171 F-NTZRP1000171   2468     NTZRP1000202 F-NTZRP1000191   2469     NTZRP1000220 F-NTZRP1000220   2470     NTZRP1000221 F-NTZRP1000220   2472     NTZRP1000324 F-NTZRP1000234   2471     NTZRP1000324 F-NTZRP1000234   2474     NTZRP1000325 F-NTZRP1000326   2475     NTZRP1000337 F-NTZRP1000348   2477     NTZRP1000348 F-NTZRP1000348   2477     NTZRP1000348 F-NTZRP1000348   2477     NTZRP1000348 F-NTZRP1000348   2477     NTZRP1000348 F-NTZRP1000348   2478     NTZRP1000356 F-NTZRP1000350   2480     NTZRP1000360 F-NTZRP1000360   2481     NTZRP1000416 F-NTZRP1000416   2484     NTZRP1000418 F-NTZRP1000418   2485     NTZRP1000404 F-NTZRP1000418   2485     NTZRP1000404 F-NTZRP1000418   2485     NTZRP1000405 F-NTZRP1000418   2485     NTZRP1000406 F-NTZRP1000418   2486     NTZRP1000407 F-NTZRP1000418   2486     NTZRP1000408 F-NTZRP1000418   2486     NTZRP1000409 F-NTZRP1000418   2490     NTZRP1000418 F-NTZRP1000418   2490     NTZRP1000418 F-NTZRP1000418   2490     NTZRP1000418 F-NTZRP1000418   2490     NTZRP1000577 F-NTZRP1000577   2497     NTZRP1000577 F-NTZRP1000577   2497     NTZRP1000581 F-NTZRP1000581   2498     NTZRP1000581 F-NTZRP1000581   2498     NTZRP1000581 F-NTZRP1000581   2498     NTZRP1000581 F-N		NT2RM4002623	F-NT2RM4002623	2455	R-NT2RM4002623	7671
NT2RP1000040		NT2RP1000018	F-NT2RP1000018	2456		
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NT2RP1000086		NT2RP1000040	F-NT2RP1000040	2458		
NTZRP1000101	•	NT2RP1000063	F-NT2RP1000063	2459		
NTZRP1000101		NT2RP1000086	F-NT2RP1000086	2460		
NT2RP1000111	10		F-NT2RP1000101	2461		
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NT2RP1000130			F-NT2RP1000112	2463		
NT2RP1000163		NT2RP1000124	F-NT2RP1000124	2464		
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NT2RP1000202	22	NT2RP1000191	F-NT2RP1000191	2469		
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	NT2RP2002208	F-NT2RP2002208	2778	R-NT2RP2002208	7859
5	NT2RP2002219	F-NT2RP2002219	2779	R-NT2RP2002219	7860
	NT2RP2002231	F-NT2RP2002231	2780	R-NT2RP2002231	7861
	NT2RP2002235	F-NT2RP2002235	2781		
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10	NT2RP2002256	F-NT2RP2002256	2783	R-NT2RP2002256	7863
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	NT2RP2002292	F-NT2RP2002292	2786	R-NT2RP2002292	7866
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	NT2RP2002333	F-NT2RP2002333	2790	R-NT2RP2002333	7870
	NT2RP2002373	F-NT2RP2002373	2791		
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	NT2RP2002394	F-NT2RP2002394	2793	R-NT2RP2002394	7872
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	NT2RP2002426	F-NT2RP2002426	2795	R-NT2RP2002426	7874
25	NT2RP2002439	F-NT2RP2002439	2796	R-NT2RP2002439	7875
	NT2RP2002442	F-NT2RP2002442	2797		
	NT2RP2002457	F-NT2RP2002457	2798	R-NT2RP2002457	7876
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10	NT2RP 2002800	F-NT2RP2002800	2829	R-NT2RP2002800	7907
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	NT2RP2002862	F-NT2RP2002862	2832	R-NT2RP2002862	7910
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	NT2RP2002929	F-NT2RP2002929	2837	R-NT2RP2002929	7915
20	NT2RP2002939	F-NT2RP2002939	2838		
	NT2RP2002954	F-NT2RP2002954	2839	R-NT2RP2002954	7916
	NT2RP2002959	F-NT2RP2002959	2840	R-NT2RP2002959	7917
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25	NT2RP2002980	F-NT2RP2002980	2842	R-NT2RP2002980	7919
	NT2RP2002986	F-NT2RP2002986	2843	R-NT2RP2002986	7920
	NT2RP2002987	F-NT2RP2002987	2844	R-NT2RP2002987	7921
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	NT2RP2003073	F-NT2RP2003073	2848	R-NT2RP2003073	7925
	NT2RP2003099	F-NT2RP2003099	2849	R-NT2RP2003099	7926
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	NT2RP2003117	F-NT2RP2003117	2851	R-NT2RP2003117	7928
	NT2RP2003121	F-NT2RP2003121	2852	R-NT2RP2003121	7929
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	NT2RP2003157	F-NT2RP2003157	2856		
	NT2RP2003158	F-NT2RP2003158	2857		2000
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	NT2RP2003293	F-NT2RP2003293	2873	R-NT2RP2003293	7947
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	NT2RP2003543	F-NT2RP2003543	2899	R-NT2RP2003543	7971
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	NT2RP2003567	F-NT2RP2003567	2902		
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	NT2RP2003713	F-NT2RP2003713	2914	R-NT2RP2003713	7985
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15	NT2RP2003825	F-NT2RP2003825	2926		
	NT2RP2003840	F-NT2RP2003840	2927	R-NT2RP2003840	7997
	NT2RP2003857	F-NT2RP2003857	2928	R-NT2RP2003857	7998
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	NT2RP2003981	F-NT2RP2003981	2936	R-NT2RP2003981	8006
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70	NT2RP2004124	F-NT2RP2004124	2947	R-NT2RP2004124	8016
	NT2RP2004142	F-NT2RP2004142	2948	R-NT2RP2004142	8017
	NT2RP2004152	F-NT2RP2004152	2949	R-NT2RP2004152	8018
	NT2RP2004165	F-NT2RP2004165	2950	R-NT2RP2004165	8019
45	NT2RP2004170	F-NT2RP2004170	2951	R-NT2RP2004170	8020
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	NT2RP2004226	F-NT2RP2004226	2957	R-NT2RP2004226	8026
	NT2RP2004232	F-NT2RP2004232	2958	R-NT2RP2004232	8027
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	NT2RP2004242	F-NT2RP2004242	2961	R-NT2RP2004242	8030
	NT2RP2004245	F-NT2RP2004245	2962	R-NT2RP2004245	8031
5	NT2RP2004270	F-NT2RP2004270	2963	R-NT2RP2004270	8032
	NT2RP2004300	F-NT2RP2004300	2964	R-NT2RP2004300	8033
	NT2RP2004316	F-NT2RP2004316	2965	R-NT2RP2004316	8034
	NT2RP2004321	F-NT2RP2004321	2966	R-NT2RP2004321	8035
10	NT2RP2004339	F-NT2RP2004339	2967	R-NT2RP2004339	8036
.•	NT2RP2004347	F-NT2RP2004347	2968	R-NT2RP2004347	8037
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	NT2RP2004365	F-NT2RP2004365	2970	R-NT2RP2004365	8039
15	NT2RP2004366	F-NT2RP2004366	2971	R-NT2RP2004366	8040
15	NT2RP2004373	F-NT2RP2004373	2972	R-NT2RP2004373	8041
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25	NT2RP2004463	F-NT2RP2004463	2980		
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	NT2RP2004568	F-NT2RP2004568	2987	R-NT2RP2004568	8055
35	NT2RP2004580	F-NT2RP2004580	2988	R-NT2RP2004580	8056
	NT2RP2004587	F-NT2RP2004587	2989	R-NT2RP2004587	8057
	NT2RP2004594	F-NT2RP2004594	2990	R-NT2RP2004594	8058
	NT2RP2004600	F-NT2RP2004600	2991	R-NT2RP2004600	8059
40	NT2RP2004602	F-NT2RP2004602	2992	R-NT2RP2004602	8060
40	NT2RP2004614	F-NT2RP2004614	2993	R-NT2RP2004614	8061
	NT2RP2004655	F-NT2RP2004655	2994	R-NT2RP2004655	8062
	NT2RP2004664		2995	R-NT2RP2004664	8063
	NT2RP2004675		2996	R-NT2RP2004675	8064
45	NT2RP2004681	F-NT2RP2004681	2997	R-NT2RP2004681	8065
	NT2RP2004689	F-NT2RP2004689	2998	R-NT2RP2004689	8066
	NT2RP2004709	F-NT2RP2004709	2999	R-NT2RP2004709	8067
	NT2RP2004710	F-NT2RP2004710	3000	R-NT2RP2004710	8068
50	NT2RP2004736		3001	R-NT2RP2004736	8069
	NT2RP2004743		3002	R-NT2RP2004743	8070
	NT2RP2004767		3003	R-NT2RP2004767	8071
	NT2RP2004768		3004		0000
55	NT2RP2004775	F-NT2RP2004775	3005	R-NT2RP2004775	8072

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		NT2RP2004799	F-NT2RP2004799	3007	R-NT2RP2004799	8074
		NT2RP2004802	F-NT2RP2004802	3008	R-NT2RP2004802	8075
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		NT2RP2004841	F-NT2RP2004841	3010	R-NT2RP2004841	8077
		NT2RP2004861	F-NT2RP2004861	3011	R-NT2RP2004861	8078
		NT2RP2004897	F-NT2RP2004897	3012	R-NT2RP2004897	8079
10		NT2RP2004933	F-NT2RP2004933	3013		
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		NT2RP2004959	F-NT2RP2004959	3015	R-nnnnnnnnnnn	8081
		NT2RP2004961	F-NT2RP2004961	3016	R-NT2RP2004961	8082
		NT2RP2004962	F-NT2RP2004962	3017	R-NT2RP2004962	8083
15		NT2RP2004967	F-NT2RP2004967	3018	R-NT2RP2004967	8084
		NT2RP2004978	F-NT2RP2004978	3019	R-NT2RP2004978	8085
		NT2RP2004982	F-NT2RP2004982	3020	R-NT2RP2004982	8086
		NT2RP2004985	F-NT2RP2004985	3021	R-NT2RP2004985	8087
20	1	NT2RP2004999	F-NT2RP2004999	3022	R-NT2RP2004999	8088
		NT2RP2005000	F-NT2RP2005000	3023	R-NT2RP2005000	8089
		NT2RP2005001	F-NT2RP2005001	3024	R-NT2RP2005001	8090
		NT2RP2005003	F-NT2RP2005003	3025	R-NT2RP2005003	8091
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		NT2RP2005020	F-NT2RP2005020	3028	R-NT2RP2005020	8094
		NT2RP2005022	F-NT2RP2005022	3029		
30	)	NT2RP2005031	F-NT2RP2005031	3030	R-NT2RP2005031	8095
		NT2RP2005037	F-NT2RP2005037	3031	R-NT2RP2005037	8096
		NT2RP2005038	F-NT2RP2005038	3032	R-NT2RP2005038	8097
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50	NT2RP3001004	F-NT2RP3001004	3277	R-NT2RP3001004	8331
	NT2RP3001007	F-NT2RP3001007	3278	R-NT2RP3001007	8332
	NT2RP3001055	F-NT2RP3001055	3279	R-NT2RP3001055	8333
	NT2RP3001057	F-NT2RP3001057	3280	R-NT2RP3001057	8334
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15	NT2RP3001140	F-NT2RP3001140	3294	R-NT2RP3001140	8348
	NT2RP3001147	F-NT2RP3001147,	3295	R-NT2RP3001147	8349
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	NT2RP3001214	F-NT2RP3001214	3299	R-NT2RP3001214	8353
	NT2RP3001216	F-NT2RP3001216	3300	R-NT2RP3001216	8354
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25	NT2RP3001232	F-NT2RP3001232	3302	R-NT2RP3001232	8356
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	NT2RP3001239	F-NT2RP3001239	3304	R-NT2RP3001239	8358
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	NT2RP3001819	F-NT2RP3001819	3379	R-NT2RP3001819	8429
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25	NT2RP3001989	F-NT2RP3001989	3394	R-NT2RP3001989	8443
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35	NT2RP3002056	F-NT2RP3002056	3402	R-NT2RP3002056	8451
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	NT2RP3002062	F-NT2RP3002062	3404	R-NT2RP3002062	8453
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40	NT2RP3002081	F-NT2RP3002081	3406	R-NT2RP3002081	8455
40	NT2RP3002097	F-NT2RP3002097	3407	R-NT2RP3002097	8456
	NT2RP3002102	F-NT2RP3002102	3408	R-NT2RP3002102	8457
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45	NT2RP3002146	F-NT2RP3002146	3411	R-NT2RP3002146	8459
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	NT2RP3002151	F-NT2RP3002151	3413	R-NT2RP3002151	8461
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50	NT2RP3002165	F-NT2RP3002165	3415	R-NT2RP3002165	8463
	NT2RP3002166	F-NT2RP3002166	3416	R-NT2RP3002166	8464
	NT2RP3002173	F-NT2RP3002173	3417	R-NT2RP3002173	8465
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55	NT2RP3002244	F-NT2RP3002244	3419	R-NT2RP3002244	8467

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	NT2RP3002303	F-NT2RP3002303	3424	R-NT2RP3002303	8472
	NT2RP3002304	F-NT2RP3002304	3425	R-NT2RP3002304	8473
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10	NT2RP3002343	F-NT2RP3002343	3427	R-NT2RP3002343	8475
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	NT2RP3002352	F-NT2RP3002352	3429	R-NT2RP3002352	8477
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	NT2RP3002455	F-NT2RP3002455,	3433	R-NT2RP3002455	8478
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25	NT2RP3002566	F-NT2RP3002566	3440	R-NT2RP3002566	8485
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40	NT2RP3002682	F-NT2RP3002682	3452	R-NT2RP3002682	8495
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	NT2RP3002861	F-NT2RP3002861	3463	R-NT2RP3002861	8506
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	NT2RP3003204	F-NT2RP3003204	3496	R-NT2RP3003204	8537
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45	NT2RP3003278	F-NT2RP3003278	3503	R-NT2RP3003278	8543
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	NT2RP3003290	F-NT2RP3003290	3505	R-NT2RP3003290	8545
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50	NT2RP3003302	F-NT2RP3003302	3507	R-NT2RP3003302	8547
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	NT2RP3003491	F-NT2RP3003491,	3525	R-NT2RP3003491	8565
	NT2RP3003500	F-NT2RP3003500	3526	R-NT2RP3003500	8566
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	NT2RP3003572	F-NT2RP3003572	3531	R-NT2RP3003572	8571
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	NT2RP3003589	F-NT2RP3003589	3533	R-NT2RP3003589	8573
	NT2RP3003621	F-NT2RP3003621	3534		
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	NT2RP3003656	F-NT2RP3003656	3536	R-NT2RP3003656	8575
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	NT2RP3003665	F-NT2RP3003665	3538	R-NT2RP3003665	8577
	NT2RP3003672	F-NT2RP3003672	3539	R-NT2RP3003672	8578
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	NT2RP3003842	F-NT2RP3003842	3556	R-NT2RP3003842	8594
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		NT2RP3004282	F-NT2RP3004282	3588		
		NT2RP3004332	F-NT2RP3004332	3589	D \mannaa	
40		NT2RP3004334	F-NT2RP3004334	3590	R-NT2RP3004334	8625
40		NT2RP3004341	F-NT2RP3004341	3591	R-NT2RP3004341	8626
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	NT2RP3004507	F-NT2RP3004507	3609	R-NT2RP3004507	8644
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50	NT2RP4000259	F-NT2RP4000259	3645	R-NT2RP4000259	8677
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	NT2RP4000781	F-NT2RP4000781	3689	R-NT2RP4000781	8716
	NT2RP4000787	F-NT2RP4000787	3690		
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	NT2RP4000839		3694		
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	NT2RP4000879	F-NT2RP4000879	3698	R-NT2RP4000879	8723
5	NT2RP4000907	F-NT2RP4000907	3699	R-nnnnnnnnnnn	8724
	NT2RP4000915	F-NT2RP4000915	3700	R-nnnnnnnnnnn	8725
	NT2RP4000918	F-NT2RP4000918	3701		
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	NT2RP4000929	F-NT2RP4000929	3705	R-NT2RP4000929	8729
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15	NT2RP4000973	F-NT2RP4000973	3707	R-NT2RP4000973	8731
	NT2RP4000975	F-NT2RP4000975	3708	R-NT2RP4000975	8732
	NT2RP4000979	F-NT2RP4000979,	3709	R-NT2RP4000979	8733
	NT2RP4000984	F-NT2RP4000984	3710	R-NT2RP4000984	8734
	NT2RP4000989	F-NT2RP4000989	3711	R-NT2RP4000989	8735
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	NT2RP4000997	F-NT2RP4000997	3713	R-NT2RP4000997	8737
	NT2RP4001004	F-NT2RP4001004	3714	R-NT2RP4001004	8738
	NT2RP4001006	F-NT2RP4001006	3715	R-NT2RP4001006	8739
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	NT2RP4001041	F-NT2RP4001041	3718	R-NT2RP4001041	8742
	NT2RP4001057	F-NT2RP4001057	3719	R-NT2RP4001057	8743
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	NT2RP4001079	F-NT2RP4001079	3722	R-NT2RP4001079	8746
	NT2RP4001080	F-NT2RP4001080	3723	R-NT2RP4001080	8747
35	NT2RP4001086	F-NT2RP4001086	3724	R-nnnnnnnnnnnn	8748
	NT2RP4001095	F-NT2RP4001095	3725	R-NT2RP4001095	8749
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40	NT2RP4001122	F-NT2RP4001122	3728	R-NT2RP4001122	8752
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	NT2RP4001143	F-NT2RP4001143	3731	R-NT2RP4001143	8755
45	NT2RP4001148	F-NT2RP4001148	3732	R-NT2RP4001148	8756
45	NT2RP4001149	F-NT2RP4001149	3733	R-NT2RP4001149	8757
	NT2RP4001150	F-NT2RP4001150	3734	R-NT2RP4001150	8758
	NT2RP4001159	F-NT2RP4001159	3735	R-NT2RP4001159	8759
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50	NT2RP4001206	F-NT2RP4001206	3737	R-nnnnnnnnnn	8761
	NT2RP4001207	F-NT2RP4001207	3738	R-NT2RP4001207	8762
	NT2RP4001210	F-NT2RP4001210	3739	R-NT2RP4001210	8763
	NT2RP4001213	F-NT2RP4001213	3740	R-NT2RP4001213	8764
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	NT2RP4001442	F-NT2RP4001442	3764	R-NT2RP4001442	8786
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30	NT2RP4001474	F-NT2RP4001474	3766	R-NT2RP4001474	8788
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25	NT2RP4001507	F-NT2RP4001507	3770	R-NT2RP4001507	8792
35	NT2RP4001524	F-NT2RP4001524	3771	R-NT2RP4001524	8793
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	NT2RP4001638	F-NT2RP4001638	3785	R-NT2RP4001638	8807
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	NT2RP4001679	F-NT2RP4001679	3789		
	NT2RP4001696	F-NT2RP4001696	3790	R-NT2RP4001696	8811
5	NT2RP4001725	F-NT2RP4001725	3791	R-NT2RP4001725	8812
	NT2RP4001730	F-NT2RP4001730	3792	R-nnnnnnnnnn	8813
	NT2RP4001739	F-NT2RP4001739	3793	R-NT2RP4001739	8814
	NT2RP4001753	F-NT2RP4001753	3794	R-NT2RP4001753	8815
10	NT2RP4001760	F-NT2RP4001760	3795	R-NT2RP4001760	8816
10	NT2RP4001790	F-NT2RP4001790	3796	R-NT2RP4001790	8817
	NT2RP4001803	F-NT2RP4001803	3797	R-NT2RP4001803	8818
	NT2RP4001822	F-NT2RP4001822	3798	R-NT2RP4001822	8819
	NT2RP4001823	F-NT2RP4001823	3799	R-NT2RP4001823	8820
15	NT2RP4001828	F-NT2RP4001828	3800	R-NT2RP4001828	8821
	NT2RP4001838	F-NT2RP4001838,	3801	R-NT2RP4001838	8822
	NT2RP4001841	F-NT2RP4001841	3802		
	NT2RP4001849	F-NT2RP4001849	3803	R-NT2RP4001849	8823
20	NT2RP4001861	F-NT2RP4001861	3804		
	NT2RP4001889	F-NT2RP4001889	3805	R-NT2RP4001889	8824
	NT2RP4001893	F-NT2RP4001893	3806	R-NT2RP4001893	8825
	NT2RP4001896	F-NT2RP4001896	3807	R-NT2RP4001896	8826
25	NT2RP4001901	F-NT2RP4001901	3808	R-NT2RP4001901	8827
	NT2RP4001927	F-NT2RP4001927	3809	R-NT2RP4001927	8828
	NT2RP4001938	F-NT2RP4001938	3810	R-NT2RP4001938	8829
	NT2RP4001946	F-NT2RP4001946	3811	R-NT2RP4001946	8830
30	NT2RP4001950	F-NT2RP4001950	3812	R-NT2RP4001950	8831
	NT2RP4001953	F-NT2RP4001953	3813	R-NT2RP4001953	8832
	NT2RP4001966	F-NT2RP4001966	3814	R-NT2RP4001966	8833
	NT2RP4001975	F-NT2RP4001975	3815	R-NT2RP4001975	8834
35	NT2RP4002018	F-NT2RP4002018	3816	R-NT2RP4002018	8835
	NT2RP4002047	F-NT2RP4002047	3817	R-NT2RP4002047	8836
	NT2RP4002052	F-NT2RP4002052	3818	R-NT2RP4002052	8837
	NT2RP4002058	F-NT2RP4002058	3819	R-NT2RP4002058	8838
40	NT2RP4002071	F-NT2RP4002071	3820	R-NT2RP4002071	8839
40	NT2RP4002075	F-NT2RP4002075	3821	R-NT2RP4002075	8840
	NT2RP4002078	F-NT2RP4002078	3822	R-NT2RP4002078	8841
	NT2RP4002081	F-NT2RP4002081	3823	R-nnnnnnnnnnn	8842
	NT2RP4002083	F-NT2RP4002083	3824	R-NT2RP4002083	8843
45	NT2RP4002408	F-NT2RP4002408	3825	R-NT2RP4002408	8844
	NT2RP4002791	F-NT2RP4002791	3826	R-NT2RP4002791	8845
	NT2RP4002888	F-NT2RP4002888	3827	R-NT2RP4002888	8846
	NT2RP4002905	F-NT2RP4002905	3828	R-NT2RP4002905	8847
50	NT2RP5003459	F-NT2RP5003459	3829	•	
	NT2RP5003461	F-NT2RP5003461	3830		
	NT2RP5003477	F-NT2RP5003477	3831		
	NT2RP5003492	F-NT2RP5003492	3832		
55	NT2RP5003500	F-NT2RP5003500	3833		

	NT2RP5003506	F-NT2RP5003506	3834		
	NT2RP5003512	F-NT2RP5003512	3835		
	NT2RP5003522	F-NT2RP5003522	3836		
5	NT2RP5003524	F-NT2RP5003524	3837		
	NT2RP5003534	F-NT2RP5003534	3838		
	0VARC1000001	F-0VARC1000001	3839	R-0VARC1000001	8848
	0VARC1000004	F-0VARC1000004	3840	R-0VARC1000004	8849
10	OVARC1000006	F-0VARC1000006	3841	R-0VARC1000006	8850
	0VARC1000013	F-0VARC1000013	3842	R-0VARC1000013	8851
	OVARC1000014	F-0VARC1000014	3843	R-0VARC1000014	8852
	OVARC1000017	F-0VARC1000017	3844	R-OVARC1000017	8853
15	0VARC1000035	F-0VARC1000035	3845	R-0VARC1000035	8854
13	OVARC1000058	F-0VARC1000058	3846	R-0VARC1000058	8855
	OVARC1000060	F-0VARC1000060,	3847	R-0VARC1000060	8856
	0VARC1000068	F-0VARC1000068	3848	R-0VARC1000068	8857
	OVARC1000071	F-0VARC1000071	3849	R-OVARC1000071	8858
20	OVARC1000085	F-0VARC1000085	3850	R-OVARC1000085	8859
	0VARC1000087	F-0VARC1000087	3851	R-nnnnnnnnnnn	8860
	0VARC1000091	F-0VARC1000091	3852	R-0VARC1000091	8861
•	OVARC1000092	F-0VARC1000092	3853	R-0VARC1000092	8862
25	OVARC1000106	F-0VARC1000106	3854	R-OVARC1000106	8863
	OVARC1000109	F-0VARC1000109	3855		
	OVARC1000113	F-0VARC1000113	3856	R-0VARC1000113	8864
	OVARC1000114	F-0VARC1000114	3857	R-0VARC1000114	8865
30	OVARC1000133	F-0VARC1000133	3858	R-OVARC1000133	8866
	0VARC1000139	F-0VARC1000139	3859		
	0VARC1000145	F-0VARC1000145	3860	R-OVARC1000145	8867
	0VARC1000148	F-OVARC1000148	3861	R-OVARC1000148	8868
35	OVARC1000151	F-0VARC1000151	3862	R-0VARC1000151	8869
	0VARC1000168	F-0VARC1000168	3863	R-0VARC1000168	8870
	OVARC1000191	F-0VARC1000191	3864	R-0VARC1000191	8871
·	0VARC1000198	F-0VARC1000198	3865	R-0VARC1000198	8872
40	OVARC1000209	F-0VARC1000209	3866	R-0VARC1000209	8873
40	OVARC1000212	F-0VARC1000212	3867	R-OVARC1000212	8874
	0VARC1000240	F-0VARC1000240	3868	R-0VARC1000240	8875
	OVARC1000241	F-0VARC1000241	3869	R-OVARC1000241	8876
	0VARC1000288	F-0VARC1000288	3870	R-0VARC1000288	8877
45	OVARC1000302	F-0VARC1000302	3871	R-0VARC1000302	8878
	0VARC1000304	F-0VARC1000304	3872	R-0VARC1000304	8879
	0VARC1000309	F-0VARC1000309	3873	R-0VARC1000309	8880
	OVARC1000321	F-0VARC1000321	3874	R-0VARC1000321	8881
50	0VARC1000326	F-0VARC1000326	3875	R-0VARC1000326	8882
	0VARC1000335	F-0VARC1000335	3876	R-0VARC1000335	8883
	0VARC1000347	F-0VARC1000347	3877	R-OVARC1000347	8884
	0VARC1000384	F-0VARC1000384	3878	R-0VARC1000384	8885
55	0VARC1000408	F-0VARC1000408	3879	R-0VARC1000408	8886

	0VARC1000411	F-0VARC1000411	3880	R-OVARC1000411	8887
	OVARC1000414	F-0VARC1000414	3881	R-OVARC1000414	8888
	OVARC1000420	F-0VARC1000420	3882	R-OVARC1000420	8889
5	OVARC1000427	F-0VARC1000427	3883	R-OVARC1000427	8890
	0VARC1000431	F-0VARC1000431	3884	R-OVARC1000431	8891
	0VARC1000437	F-0VARC1000437	3885	R-OVARC1000437	8892
	OVARC1000440	F-0VARC1000440	3886	R-OVARC1000440	8893
10	OVARC1000442	F-0VARC1000442	3887	R-OVARC1000442	8894
	0VARC1000443	F-0VARC1000443	3888	R-OVARC1000443	8895
	OVARC1000461	F-0VARC1000461	3889	R-OVARC1000461	8896
	OVARC1000465	F-0VARC1000465	3890	R-OVARC1000465	8897
15	0VARC1000466	F-0VARC1000466	3891	R-OVARC1000466	8898
10	0VARC1000473	F-0VARC1000473	3892	R-0VARC1000473	8899
	0VARC1000479	F-0VARC1000479,	3893	R-OVARC1000479	8900
	0VARC1000486	F-0VARC1000486	3894	R-0VARC1000486	8901
	0VARC1000496	F-0VARC1000496	3895	R-0VARC1000496	8902
20	0VARC1000520	F-0VARC1000520	3896	R-OVARC1000520	8903
	0VARC1000526	F-0VARC1000526	3897	R-OVARC1000526	8904
	0VARC1000533	F-0VARC1000533	3898	R-0VARC1000533	8905
	0VARC1000543	F-0VARC1000543	3899	R-OVARC1000543	8906
25	0VARC1000556	F-0VARC1000556	3900	R-0VARC1000556	8907
	OVARC1000557	F-0VARC1000557	3901	R-OVARC1000557	8908
	OVARC1000564	F-0VARC1000564	3902	R-OVARC1000564	8909
	0VARC1000573	F-0VARC1000573	3903	R-OVARC1000573	8910
30	OVARC1000576	F-0VARC1000576	3904		
	0VARC1000578	F-0VARC1000578	3905	R-0VARC1000578	8911
	0VARC1000588	F-0VARC1000588	3906	R-OVARC1000588	8912
	0VARC1000605	F-0VARC1000605	3907	R-0VARC1000605	8913
35	0VARC1000622	F-0VARC1000622	3908	R-OVARC1000622	8914
	OVARC1000640	F-0VARC1000640	3909	R-0VARC1000640	8915
	0VARC1000649	F-0VARC1000649	3910		
	0VARC1000661	F-0VARC1000661	3911	R-0VARC1000661	8916
40	0VARC1000678	F-0VARC1000678	3912	R-OVARC1000678	8917
	0VARC1000679	F-0VARC1000679	3913	R-nnnnnnnnnnn	8918
	0VARC1000681	F-0VARC1000681	3914	R-OVARC1000681	8919
	0VARC1000682	F-0VARC1000682	3915		
45	0VARC1000689	F-0VARC1000689	3916	R-OVARC1000689	8920
45	OVARC1000700	F-0VARC1000700	3917	R-0VARC1000700	8921
•	OVARC1000703	F-0VARC1000703	3918	R-OVARC1000703	8922
	OVARC1000722	F-0VARC1000722	3919		
	OVARC1000730	F-0VARC1000730	3920	R-OVARC1000730	8923
50	OVARC1000746	F-0VARC1000746	3921	R-OVARC1000746	8924
	OVARC1000769	F-0VARC1000769	3922	R-OVARC1000769	8925
	0VARC1000771	•	3923	R-0VARC1000771	8926
	0VARC1000781	F-0VARC1000781	3924	R-0VARC1000781	8927
55	OVARC1000787	F-0VARC1000787	3925	R-OVARC1000787	8928

	OVARC1000800	F-0VARC1000800	3926	R-0VARC1000800	8929
	0VARC1000802	F-0VARC1000802	3927	R-0VARC1000802	8930
	OVARC1000834	F-0VARC1000834	3928	R-0VARC1000834	8931
5	OVARC1000846	F-0VARC1000846	3929	R-0VARC1000846	8932
	OVARC1000850	F-0VARC1000850	3930	R-0VARC1000850	8933
	OVARC1000862	F-0VARC1000862	3931	R-0VARC1000862	8934
	OVARC1000876	F-0VARC1000876	3932	R-0VARC1000876	8935
10	OVARC1000883	F-0VARC1000883	3933	R-0VARC1000883	8936
	OVARC1000885	F-0VARC1000885	3934	R-0VARC1000885	8937
	OVARC1000886	F-0VARC1000886	3935	R-0VARC1000886	8938
	OVARC1000890	F-0VARC1000890	3936		
15	0VARC1000891	F-0VARC1000891	3937	R-0VARC1000891	8939
15	0VARC1000897	F-0VARC1000897	3938	R-0VARC1000897	8940
	OVARC1000912	F-0VARC1000912,	3939	R-0VARC1000912	8941
	OVARC1000915	F-0VARC1000915	3940	R-0VARC1000915	8942
	OVARC1000924	F-0VARC1000924	3941	R-0VARC1000924	8943
20	0VARC1000936	F-0VARC1000936	3942	R-0VARC1000936	8944
	OVARC1000937	F-0VARC1000937	3943	R-0VARC1000937	8945
	OVARC1000945	F-0VARC1000945	3944	R-0VARC1000945	8946
	OVARC1000948	F-0VARC1000948	3945	R-0VARC1000948	8947
25	OVARC1000959	F-0VARC1000959	3946	R-0VARC1000959	8948
	OVARC1000960	F-0VARC1000960	3947	R-0VARC1000960	8949
	OVARC1000964	F-0VARC1000964	3948		
	OVARC1000971	F-0VARC1000971	3949	R-0VARC1000971	8950
30	OVARC1000984	F-0VARC1000984	3950	R-0VARC1000984	8951
	OVARC1000996	F-0VARC1000996	3951	R-0VARC1000996	8952
	OVARC1000999	F-0VARC1000999	3952	R-0VARC1000999	8953
	OVARC1001000	F-0VARC1001000	3953	R-0VARC1001000	8954
35	OVARC1001004	F-0VARC1001004	3954	R-0VARC1001004	8955
	OVARC1001010	F-0VARC1001010	3955	R-0VARC1001010	8956
	OVARC1001011	F-0VARC1001011	3956	R-0VARC1001011	8957
	OVARC1001032	F-0VARC1001032	3957	R-0VARC1001032	8958
40	0VARC1001034	F-0VARC1001034	3958	R-0VARC1001034	8959
40	0VARC1001038	F-0VARC1001038	3959	R-0VARC1001038	8960
	0VARC1001040	F-0VARC1001040	3960	R-0VARC1001040	8961
	OVARC1001044	F-0VARC1001044	3961	R-0VARC1001044	8962
	OVARC1001051	F-0VARC1001051	3962	R-0VARC1001051	8963
45	OVARC1001055	F-0VARC1001055	3963	R-0VARC1001055	8964
	OVARC1001062	F-0VARC1001062	3964	R-0VARC1001062	8965
	OVARC1001065	F-0VARC1001065	3965		
	OVARC1001068	F-0VARC1001068	3966	R-0VARC1001068	8966
50	OVARC1001072	F-0VARC1001072	3967	R-0VARC1001072	8967
	OVARC1001074	F-0VARC1001074	3968	R-0VARC1001074	8968
	OVARC1001085	F-0VARC1001085	3969	R-0VARC1001085	8969
	OVARC1001092	F-0VARC1001092	3970	R-0VARC1001092	8970
55	OVARC1001107	F-0VARC1001107	3971		

	OVARC1001113	F-0VARC1001113	3972	R-OVARC1001113	8971
	OVARC1001117	F-0VARC1001117	3973	R-0VARC10011117	8972
	0VARC1001118	F-0VARC1001118	3974	R-0VARC1001118	8973
5	0VARC1001129	F-0VARC1001129	3975	R-0VARC1001129	8974
	OVARC1001154	F-0VARC1001154	3976		
	OVARC1001161	F-0VARC1001161	3977	R-0VARC1001161	8975
	OVARC1001162	F-0VARC1001162	3978	R-0VARC1001162	8976
1Ő	OVARC1001167	F-0VARC1001167	3979	R-0VARC1001167	8977
	OVARC1001169	F-0VARC1001169	3980	R-OVARC1001169	8978
	OVARC1001170	F-0VARC1001170	3981	R-0VARC1001170	8979
	OVARC1001171	F-0VARC1001171	3982		
15	OVARC1001173	F-0VARC1001173	3983	R-0VARC1001173	8980
,,	OVARC1001176	F-0VARC1001176	3984		
	OVARC1001180	F-0VARC1001180,	3985	R-0VARC1001180	8981
	OVARC1001188	F-0VARC1001188	3986	R-0VARC1001188	8982
	OVARC1001200	F-0VARC1001200	3987	R-0VARC1001200	8983
20	0VARC1001232	F-0VARC1001232	3988	R-0VARC1001232	8984
	OVARC1001240	F-0VARC1001240	3989	R-0VARC1001240	8985
	0VARC1001243	F-0VARC1001243	3990	R-0VARC1001243	8986
	0VARC1001244	F-0VARC1001244	3991		
25	0VARC1001261	F-0VARC1001261	3992	R-0VARC1001261	8987
	0VARC1001268	F-0VARC1001268	3993	R-0VARC1001268	8988
	0VARC1001270	F-0VARC1001270	3994	R-0VARC1001270	8989
	OVARC1001271	F-0VARC1001271	3995	R-0VARC1001271	8990
30	OVARC1001282	F-0VARC1001282	3996	R-0VARC1001282	8991
	0VARC1001296	F-0VARC1001296	3997	R-0VARC1001296	8992
	0VARC1001306	F-0VARC1001306	3998	R-nannananana	8993
	OVARC1001329	F-0VARC1001329	3999	R-0VARC1001329	8994
35	0VARC1001330	F-0VARC1001330	4000	R-0VARC1001330	8995
	OVARC1001339	F-0VARC1001339	4001	R-0VARC1001339	8996
	0VARC1001341	F-0VARC1001341	4002	R-0VARC1001341	8997
	OVARC1001342	F-0VARC1001342	4003	R-0VARC1001342	8998
40	0VARC1001344	F-0VARC1001344	4004	R-0VARC1001344	8999
	OVARC1001357	F-0VARC1001357	4005	R-0VARC1001357	9000
	OVARC1001360	F-0VARC1001360	4006	R-0VARC1001360	9001
	0VARC1001369	F-0VARC1001369	4007	R-0VARC1001369	9002
45	0VARC1001372	F-0VARC1001372	4008	R-0VARC1001372	9003
45	OVARC1001376	F-0VARC1001376	4009	R-0VARC1001376	9004
	OVARC1001381	F-0VARC1001381	4010	R-0VARC1001381	9005
	OVARC1001391	F-0VARC1001391	4011	R-0VARC1001391	9006 9007
	0VARC1001399	F-0VARC1001399	4012	R-nnnnnnnnnnnn	
50	OVARC1001417 OVARC1001419	F-0VARC1001417 F-0VARC1001419	4013	R-0VARC1001417	9008
	0VARC1001419 0VARC1001425	F-0VARC1001419 F-0VARC1001425	4014	R-OVARC1001419 R-OVARC1001425	9009 9010
	0VARC1001425 0VARC1001436	F-0VARC1001425 F-0VARC1001436	4015 4016	R-0VARC1001425	9010
	OVARC1001430	F-0VARC1001442	4016	R-0VARC1001438	9011
55	01110101142	1 01111011011442	3011	W 011001001447	2012

	OVARC1001453	F-0VARC1001453	4018	R-0VARC1001453 ·	9013
	OVARC1001476	F-0VARC1001476	4019	R-0VARC1001476	9014
	OVARC1001480	F-0VARC1001480	4020	R-0VARC1001480	9015
5	OVARC1001489	F-0VARC1001489	4021	R-0VARC1001489	9016
	OVARC1001496	F-0VARC1001496	4022	R-0VARC1001496	9017
	OVARC1001506	F-0VARC1001506	4023	R-0VARC1001506	9018
	OVARC1001525	F-0VARC1001525	4024	R-OVARC1001525	9019
10	OVARC1001542	F-0VARC1001542	4025	R-0VARC1001542	9020
	OVARC1001547	F-0VARC1001547	4026	R-0VARC1001547	9021
	OVARC1001555	F-0VARC1001555	4027		
	OVARC1001577	F-0VARC1001577	4028	R-OVARC1001577	9022
15	OVARC1001600	F-0VARC1001600	4029	R-0VARC1001600	9023
13	OVARC1001610	F-0VARC1001610	4030	R-0VARC1001610	9024
	OVARC1001611	F-0VARC1001611,	4031	R-0VARC1001611	9025
	OVARC1001615	F-0VARC1001615	4032	R-OVARC1001615	9026
	OVARC1001668	F-0VARC1001668	4033	R-0VARC1001668	9027
20	OVARC1001702	F-0VARC1001702	4034	R-OVARC1001702	9028
	OVARC1001703	F-0VARC1001703	4035	R-OVARC1001703	9029
	OVARC1001711	F-0VARC1001711	4036	.R-OVARC1001711	9030
	OVARC1001713	F-0VARC1001713	4037	·	
25	OVARC1001726	F-0VARC1001726	4038	R-0VARC1001726	9031
	OVARC1001731	F-0VARC1001731	4039	R-OVARC1001731	9032
	OVARC1001745	F-0VARC1001745	4040	R-OVARC1001745	9033
	OVARC1001762	F-0VARC1001762	4041	R-nnnnnnnnnnn	9034
30	OVARC1001766	F-0VARC1001766	4042	R-0VARC1001766	9035
	OVARC1001767	F-0VARC1001767	4043	R-nnnnnnnnnnn	9036
	OVARC1001768	F-0VARC1001768	4044	R-0VARC1001768	9037
	OVARC1001791	F-0VARC1001791	4045	R-0VARC1001791	9038
35	OVARC1001795	F-0VARC1001795	4046	R-0VARC1001795	9039 9040
	OVARC1001802	F-0VARC1001802	4047	R-0VARC1001802	9040
	OVARC1001805	F-0VARC1001805	4048	R-0VARC1001805	9041
	OVARC1001809	F-0VARC1001809	4049	R-0VARC1001812	9042
40	OVARC1001812	F-0VARC1001812	4050	R-0VARC1001813	9043
40	OVARC1001813	F-0VARC1001813	4051	R-0VARC1001813	9044
	OVARC1001820	F-0VARC1001820	4052	R-0VARC1001828	9045
	OVARC1001828	F-0VARC1001828	4053	R-0VARC1001828	9046
	OVARC1001846	F-0VARC1001846	4054 4055	R-0VARC1001840	9047
45	OVARC1001861	F-0VARC1001861	4055	R-0VARC1001801	9048
	OVARC1001873	F-0VARC1001873 F-0VARC1001879	4056	R-0VARC1001879	9049
	OVARC1001879	F-0VARC1001879	4058	R-0VARC1001819	9050
	OVARC1001880 OVARC1001883		4059	R-0VARC1001883	9051
50	OVARC1001883		4060	R-0VARC1001900	9052
	OVARC1001900 OVARC1001901	-	4061	R-0VARC1001901	9053
	OVARC1001901 OVARC1001911		4062	R-0VARC1001911	9054
	OVARC1001911 OVARC1001916		4063	R-0VARC1001911	9055
55	04401001310	1 07/10/10/13/10	.000	0	

	OVARC1001928	F-0VARC1001928	4064	R-0VARC1001928	9056
	OVARC1001942	F-0VARC1001942	4065	R-0VARC1001942	9057
	OVARC1001943	F-0VARC1001943	4066	R-0VARC1001943	9058
5	OVARC1001949	F-0VARC1001949	4067	R-0VARC1001949	9059
	OVARC1001950	F-0VARC1001950	4068	R-0VARC1001950	9060
	OVARC1001987	F-0VARC1001987	4069	R-0VARC1001987	9061
	OVARC1001989	F-0VARC1001989	4070	R-0VARC1001989	9062
10	OVARC1002044	F-0VARC1002044	4071	R-0VARC1002044	9063
	OVARC1002050	F-0VARC1002050	4072	R-0VARC1002050	9064
	OVARC1002066	F-0VARC1002066	4073	R-0VARC1002066	9065
	OVARC1002082	F-0VARC1002082	4074	R-0VARC1002082	9066
4.5	OVARC1002107	F-0VARC1002107	4075	R-0VARC1002107	9067
15	OVARC1002112	F-0VARC1002112	4076		
	OVARC1002127	F-OVARC1002127,	4077	R-0VARC1002127	9068
	OVARC1002138	F-0VARC1002138	4078	R-0VARC1002138	9069
	OVARC1002143	F-OVARC1002143	4079	R-0VARC1002143	9070
20	OVARC1002156	F-0VARC1002156	4080	R-OVARC1002156	9071
	OVARC1002158	F-0VARC1002158	4081	R-0VARC1002158	9072
	OVARC1002165	F-0VARC1002165	4082	R-0VARC1002165	9073
	OVARC1002182	F-0VARC1002182	4083	R-0VARC1002182	9074
25	PLACE1000004	F-PLACE1000004	4084	R-PLACE1000004	9075
	PLACE1000005	F-PLACE1000005	4085	R-PLACE1000005	9076
	PLACE1000007	F-PLACE1000007	4086	R-PLACE1000007	9077
	PLACE1000014	F-PLACE1000014	4087	R-PLACE1000014	9078
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	PLACE1000040	F-PLACE1000040	4089	R-PLACE1000040	9080
	PLACE1000048	F-PLACE1000048	4090	R-PLACE1000048	9081
	PLACE1000050	F-PLACE1000050	4091	R-PLACE1000050	9082
35	PLACE1000061	F-PLACE1000061	4092	R-PLACE1000061	9083
55	PLACE1000066	F-PLACE1000066	4093	R-PLACE1000066	9084
	PLACE1000078	F-PLACE1000078	4094	R-PLACE1000078	9085
	PLACE1000081	F-PLACE1000081	4095	R-PLACE1000081	9086
	PLACE1000094	F-PLACE1000094	4096	R-PLACE1000094	9087
40	PLACE1000133	F-PLACE1000133	4097	R-PLACE1000133	9088
	PLACE1000142	F-PLACE1000142	4098	R-PLACE1000142	9089
	PLACE1000184	F-PLACE1000184	4099	R-PLACE1000184	9090
	PLACE1000185	F-PLACE1000185	4100	R-PLACE1000185	9091
45	PLACE1000213	F-PLACE1000213	4101	R-PLACE1000213	9092
	PLACE1000214	F-PLACE1000214	4102	R-PLACE1000214	9093
	PLACE1000236	F-PLACE1000236	4103	R-PLACE1000236	9094
	PLACE1000246	F-PLACE1000246	4104	R-PLACE1000246	9095
50	PLACE1000292	F-PLACE1000292	4105	R-PLACE1000292	9096
	PLACE1000308	F-PLACE1000308	4106		
	PLACE1000332	F-PLACE1000332	4107	R-PLACE1000332	9097
	PLACE1000347	F-PLACE1000347	4108	R-PLACE1000347	9098
55	PLACE1000374	F-PLACE1000374	4109	R-PLACE1000374	9099
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	PLACE1000380	F-PLACE1000380	4110	R-PLACE1000380	9100 -
	PLACE1000383	F-PLACE1000383	4111	R-PLACE1000383	9101
	PLACE1000401	F-PLACE1000401	4112	R-PLACE1000401	9102
5	PLACE1000406	F-PLACE1000406	4113	R-PLACE1000406	9103
	PLACE1000420	F-PLACE1000420	4114	R-PLACE1000420	9104
	PLACE1000421	F-PLACE1000421	4115	R-PLACE1000421	9105
	PLACE1000424	F-PLACE1000424	4116	R-PLACE1000424	9106
10	PLACE1000435	F-PLACE1000435	4117	R-PLACE1000435	9107
70	PLACE1000444	F-PLACE1000444	4118	R-PLACE1000444	9108
	PLACE1000453	F-PLACE1000453	4119	R-PLACE1000453	9109
	PLACE1000481	F-PLACE1000481	4120	R-PLACE1000481	9110
	PLACE1000492	F-PLACE1000492	4121	R-PLACE1000492	9111
15	PLACE1000540	F-PLACE1000540	4122	R-PLACE1000540	9112
	PLACE1000547	F-PLACE1000547,	4123	R-PLACE1000547	9113
	PLACE1000562	F-PLACE1000562	4124	R-PLACE1000562	9114
	PLACE1000564	F-PLACE1000564	4125	R-PLACE1000564	9115
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	PLACE1000588	F-PLACE1000588	4127	R-nnnnnnnnnn	9117
	PLACE1000596	F-PLACE1000596	4128	R-PLACE1000596	9118
	PLACE1000599	F-PLACE1000599	4129	R-PLACE1000599	9119
25	PLACE1000610	F-PLACE1000610	4130	R-PLACE1000610	9120
	PLACE1000611	F-PLACE1000611	4131		
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	PLACE1000653	F-PLACE1000653	4133	R-PLACE1000653	9122
30	PLACE1000656	F-PLACE1000656 ·	4134	R-PLACE1000656	9123
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	PLACE1000712	F-PLACE1000712	4136	R-PLACE1000712	9125
	PLACE1000716	F-PLACE1000716	4137	R-PLACE1000716	9126
	PLACE1000748	F-PLACE1000748	4138	R-PLACE1000748	9127
35	PLACE1000749	F-PLACE1000749	4139	R-PLACE1000749	9128
	PLACE1000755	F-PLACE1000755	4140	R-PLACE1000755	9129
	PLACE1000769	F-PLACE1000769	4141	R-PLACE1000769	9130
	PLACE1000785	F-PLACE1000785	4142	R-PLACE1000785	9131
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	PLACE1000979	F-PLACE1000979	4155	R-PLACE1000979	9144

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PLACE1001007 F-PLACE1001007 4158 R-PLACE1001007 PLACE1001010 F-PLACE1001010 4159 R-PLACE1001010 PLACE1001015 F-PLACE1001015 4160 R-PLACE1001016 PLACE1001024 F-PLACE1001024 4161 R-PLACE1001024 PLACE1001036 F-PLACE1001036 4162 R-PLACE1001036 PLACE1001054 F-PLACE1001054 4163 PLACE1001062 F-PLACE1001062 4164 R-PLACE1001065	7 9146 0 9147 5 9148 4 9149 6 9150 2 9151
PLACE1001010 F-PLACE1001010 4159 R-PLACE1001010 PLACE1001015 F-PLACE1001015 4160 R-PLACE1001010 PLACE1001024 F-PLACE1001024 4161 R-PLACE1001024 PLACE1001036 F-PLACE1001036 4162 R-PLACE1001036 PLACE1001054 F-PLACE1001054 4163 PLACE1001062 F-PLACE1001062 4164 R-PLACE1001065	5 9148 4 9149 6 9150 2 9151
PLACE1001015 F-PLACE1001015 4160 R-PLACE1001016 PLACE1001024 F-PLACE1001024 4161 R-PLACE1001024 PLACE1001036 F-PLACE1001036 4162 R-PLACE1001036 PLACE1001054 F-PLACE1001054 4163 PLACE1001062 F-PLACE1001062 4164 R-PLACE1001066	5 9148 4 9149 6 9150 2 9151
PLACE1001024 F-PLACE1001024 4161 R-PLACE1001024 PLACE1001036 F-PLACE1001036 4162 R-PLACE1001036 PLACE1001054 F-PLACE1001054 4163 PLACE1001062 F-PLACE1001062 4164 R-PLACE1001065	<ul><li>9149</li><li>9150</li><li>9151</li></ul>
PLACE1001036 F-PLACE1001036 4162 R-PLACE1001036  PLACE1001054 F-PLACE1001054 4163  PLACE1001062 F-PLACE1001062 4164 R-PLACE1001065	6 9150 2 9151
PLACE1001054 F-PLACE1001054 4163 PLACE1001062 F-PLACE1001062 4164 R-PLACE1001062	2 9151
PLACE1001062 F-PLACE1001062 4164 R-PLACE1001063	
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	J J J J J J
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PLACE 100 2007 F. D. ACE 100 1007 A 176 P. D. ACE 100 125	
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5	PLACE1001602	F-PLACE1001602	4205	R-PLACE1001602	9192
	PLACE1001603	F-PLACE1001603	4206	R-PLACE1001603	9193
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	PLACE1001632	F-PLACE1001632	4210	R-PLACE1001632	9196
	PLACE1001634	F-PLACE1001634	4211	R-PLACE1001634	9197
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	PLACE1001692	F-PLACE1001692,	4215	R-PLACE1001692	9201
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	PLACE1001716	F-PLACE1001716	4217	R-PLACE1001716	9203
20	PLACE1001720	F-PLACE1001720	4218	R-PLACE1001720	9204
	PLACE1001729	F-PLACE1001729	4219	R-PLACE1001729	9205
	PLACE1001739	F-PLACE1001739	4220	R-PLACE1001739	9206
	PLACE1001740	F-PLACE1001740	4221	R-PLACE1001740	9207
25	PLACE1001745	F-PLACE1001745	4222	R-PLACE1001745	9208
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40	PLACE1001869	F-PLACE1001869	4235	R-PLACE1001869	9219
	PLACE1001897	F-PLACE1001897	4236	R-PLACE1001897	9220
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	PLACE1001920	F-PLACE1001920	4238	R-PLACE1001920	9222
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	PLACE1001983	F-PLACE1001983	4240	R-PLACE1001983	9224
	PLACE1001989	F-PLACE1001989	4241	R-PLACE1001989	9225
	PLACE1002004	F-PLACE1002004	4242	D DI 4001000040	0000
50	PLACE1002046	F-PLACE1002046	4243	R-PLACE1002046	9226
	PLACE1002052	F-PLACE1002052	4244	R-PLACE1002052	9227
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	PLACE1002072	F-PLACE1002072	4246	R-PLACE1002072 R-PLACE1002073	9229
	PLACE1002073	F-PLACE1002073	4247	K-PLACE1002073	9230

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	PLACE1002115	F-PLACE1002115	4249	R-PLACE1002115	9232
	PLACE1002119	F-PLACE1002119	4250	R-PLACE1002119	9233
5	PLACE1002140	F-PLACE1002140	4251	R-PLACE1002140	9234
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	PLACE1002157	F-PLACE1002157	4253	R-PLACE1002157	9236
	PLACE1002163	F-PLACE1002163	4254	R-PLACE1002163	9237
10	PLACE1002170	F-PLACE1002170	4255	N I MIODITUTE	
10	PLACE1002171	F-PLACE1002171	4256	R-PLACE1002171	9238
	PLACE1002101	F-PLACE1002205	4257	R-PLACE1002205	9239
	PLACE1002213	F-PLACE1002213	4258	R-PLACE1002213	9240
	PLACE1002227	F-PLACE1002227	4259	R-PLACE1002227	9241
15	PLACE1002256	F-PLACE1002256	4260	R-PLACE1002256	9242
	PLACE1002259	F-PLACE1002259,		R-PLACE1002259	9243
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	PLACE1002319 PLACE1002342	F-PLACE1002319	4263	R-PLACE1002342	9245
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	PLACE1002395 PLACE1002399	F-PLACE1002399	4265	R-PLACE1002399	9247
		F-PLACE1002399 F-PLACE1002433	4266	R-PLACE1002333	9248
	PLACE1002433	F-PLACE1002437	4267	R-PLACE1002437	9249
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25	PLACE1002438 PLACE1002450	F-PLACE1002450	4269	R-PLACE1002450	9251
	PLACE1002450 PLACE1002465	F-PLACE1002465	4209	R-PLACE1002465	9252
	PLACE1002474	F-PLACE1002465 F-PLACE1002474	4270	R-PLACE1002474	9252
	PLACE1002474 PLACE1002477	F-PLACE1002474 F-PLACE1002477	4271	R-PLACE1002477	9254
30	PLACE1002477	F-PLACE1002477	4272	R-PLACE1002493	9255
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	PLACE1002499 PLACE1002500	F-PLACE1002499	4274	R-PLACE1002499	9257
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35	PLACE1002514 PLACE1002529	F-PLACE1002514 F-PLACE1002529	4277	R-PLACE1002514	9259
	PLACE1002525	F-PLACE1002529	4278	R-PLACE1002532	9260
	PLACE1002537	F-PLACE1002537	4279	R-PLACE1002537	9261
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40	PLACE1002578	F-PLACE1002583	4282	R-PLACE1002583	9264
	PLACE1002591	F-PLACE1002583	4283	R-PLACE1002591	9265
	PLACE1002598	F-PLACE1002591	4284	R-PLACE1002598	9266
	PLACE1002604	F-PLACE1002598	4285	R-PLACE1002604	9267
45	PLACE1002625	F-PLACE1002604 F-PLACE1002625	4286	R-PLACE1002604	9268
	PLACE1002655	F-PLACE1002655	4287	K TEACETOOZOZO	3200
	PLACE1002665	F-PLACE1002665	4288	R-PLACE1002665	9269
	PLACE1002685		4289	R-PLACE1002685	9270
50	PLACE1002685 PLACE1002714	F-PLACE1002685	4299	R-PLACE1002005	9271
	PLACE1002714 PLACE1002722	F-PLACE1002714	4290		9272
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	PLACE1002768 PLACE1002772	F-PLACE1002768 F-PLACE1002772	4292	R-PLACE1002778	9274
	1 LACE1002112	r-reaccious/12	7630	W TEMORITANS (12	2614

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	PLACE1002775	F-PLACE1002775	4294		
	PLACE1002782	F-PLACE1002782	4295	R-PLACE1002782	9275
	PLACE1002794	F-PLACE1002794	4296	R-PLACE1002794	9276
5	PLACE1002811	F-PLACE1002811	4297	R-PLACE1002811	9277
	PLACE1002815	F-PLACE1002815	4298	R-PLACE1002815	9278
	PLACE1002816	F-PLACE1002816	4299	R-PLACE1002816	9279
•	PLACE1002834	F-PLACE1002834	4300	R-PLACE1002834	9280
	PLACE1002839	F-PLACE1002839		R-PLACE1002839	9281
10			4301		
	PLACE1002851	F-PLACE1002851	4302	R-PLACE1002851	9282
	PLACE1002853	F-PLACE1002853	4303	R-PLACE1002853	9283
	PLACE1002881	F-PLACE1002881	4304	R-PLACE1002881	9284
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	PLACE1002941	F-PLACE1002941	4306	R-PLACE1002941	9286
	PLACE1002962	F-PLACE1002962,	4307	R-PLACE1002962	9287
	PLACE1002968	F-PLACE1002968	4308	R-PLACE1002968	9288
	PLACE1002991	F-PLACE1002991	4309	R-PLACE1002991	9289
20	PLACE1002993	F-PLACE1002993	4310	R-PLACE1002993	9290
	PLACE1002996	F-PLACE1002996	4311	R-PLACE1002996	9291
	PLACE1003025	F-PLACE1003025	4312	R-PLACE1003025	9292
	PLACE1003027	F-PLACE1003027	4313	R-PLACE1003027	9293
25	PLACE1003044	F-PLACE1003044	4314	R-PLACE1003044	9294
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	PLACE1003519	F-PLACE1003519	4350	R-PLACE1003519	9328
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15	PLACE1003528	F-PLACE1003528	4352	R-PLACE1003528	9330
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	PLACE1004646	F-PLACE1004646	4443	R-PLACE1004646	9419
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	PLACE1004674	F-PLACE1004674	4447	R-PLACE1004674	9423
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	PLACE1004691	F-PLACE1004691	4450	R-PLACE1004691	9426
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	PLACE1004722	F-PLACE1004722	4453	R-PLACE1004722	9429
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45	PLACE1004840	F-PLACE1004840	4469	R-PLACE1004840	9445
	PLACE1004868	F-PLACE1004868	4470	R-PLACE1004868	9446
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10	PLACE1005046	F-PLACE1005046	4486	R-PLACE1005046	9462
	PLACE1005052	F-PLACE1005052	4487	R-PLACE1005052	9463
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15	PLACE1005077	F-PLACE1005077	4490	R-PLACE1005077	9465
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	PLACE1005101	F-PLACE1005101	4493	R-PLACE1005101	9468
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25	PLACE1005146	F-PLACE1005146	4498	R-PLACE1005146	9473
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	PLACE1005261	F-PLACE1005261	4506	R-PLACE1005261	9481
35	PLACE1005266	F-PLACE1005266	4507	R-PLACE1005266	9482
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25	PLACE1005698	F-PLACE1005698	4544	R-PLACE1005698	9519
23	PLACE1005727	F-PLACE1005727	4545	R-PLACE1005727	9520
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	PLACE1005739	F-PLACE1005739	4547	R-PLACE1005739	9522
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35	PLACE1005804	F-PLACE1005804	4553	R-PLACE1005804	9528
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25	PLACE1006167	F-PLACE1006167	4590	R-PLACE1006167	9563
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	PLACE4000521	F-PLACE4000521	5233	K I LHOLIOUVISI	101.1
25	PLACE4000522	F-PLACE4000522	5234	R-PLACE4000522	10172
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	Y79AA1000037	F-Y79AA1000037	5413	-R-Y79AA1000037	10334
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30	Y79AA1000268	F-Y79AA1000268	5423	R-Y79AA1000268	10344
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	Y79AA1000410	F-Y79AA1000410	5432	R-Y79AA1000410	10353
	Y79AA1000420	F-Y79AA1C00420	5433	R-Y79AA1000420	10354
	Y79AA1000469	F-Y79AA1000469	5434	R-Y79AA1000469	10355
45	Y79AA1000480	F-Y79AA1000480	5435	R-Y79AA1000480	10356
45	Y79AA1000538	F-Y79AA1000538	5436	R-Y79AA1000538	10357
	Y79AA1000539	F-Y79AA1000539	5437	R-Y79AA1000539	10358
	Y79AA1000540	F-Y79AA1000540	5438	R-Y79AA1000540	10359
	Y79AA1000560	F-Y79AA1000560	5439	R-Y79AA1000560	10360
50	Y79AA1000574	F-Y79AA1000574	5440	R-Y79AA1000574	10361
	Y79AA1000589	F-Y79AA1000589	5441		
	Y79AA1000627	F-Y79AA1000627	5442	R-Y79AA1000627	10362
	Y79AA1000705	F-Y79AA1000705	5443	R-Y79AA1000705	10363

	Y79AA1000734	F-Y79AA1000734	5444	R-Y79AA1000734	10364
	Y79AA1000748	F-Y79AA1000748	5445	R-Y79AA1000748	10365
	Y79AA1000752	F-Y79AA1000752	5446	R-Y79AA1000752	10366
5	Y79AA1000774	F-Y79AA1000774	5447	R-Y79AA1000774	10367
	Y79AA1000782	F-Y79AA1000782	5448	R-Y79AA1000782	10368
	Y79AA1000784	F-Y79AA1000784	5449	R-Y79AA1000784	10369
	Y79AA1000794	F-Y79AA1000794	5450	R-Y79AA1000794	10370
10	Y79AA1000800	F-Y79AA1000800	5451	R-Y79AA1000800	10371
	Y79AA1000802	F-Y79AA1000802	5452	R-nnnnnnnnnnn	10372
	Y79AA1000805	F-Y79AA1000805	5453	R-Y79AA1000805	10373
	Y79AA1000824	F-Y79AA1000824	5454	R-Y79AA1000824	10374
15	Y79AA1000827	F-Y79AA1000827	5455	R-Y79AA1000827	10375
	Y79AA1000833	F-Y79AA1000833	5456		
	Y79AA1000850	F-Y79AA1000850,	5457	R-Y79AA1000850	10376
	Y79AA1000962	F-Y79AA1000962	5458	R-Y79AA1000962	10377
20	Y79AA1000966	F-Y79AA1000966	5459	•	
	Y79AA1000968	F-Y79AA1000968	5460	R-Y79AA1000968	10378
	Y79AA1000969	F-Y79AA1000969	5461	R-Y79AA1000969	10379
	Y79AA1000976	F-Y79AA1000976	5462	R-Y79AA1000976	10380
25	Y79AA1000985	F-Y79AA1000985	5463	R-Y79AA1000985	10381
23	Y79AA1001023	F-Y79AA1001023	5464	R-Y79AA1001023	10382
	Y79AA1001041	F-Y79AA1001041	5465	R-Y79AA1001041	10383
	Y79AA1001048	F-Y79AA1001048	5466	R-Y79AA1001048	10384
	Y79AA1001061	F-Y79AA1001061	5467	R-Y79AA1001061	10385
30	Y79AA1001068	F-Y79AA1001068	5468	R-Y79AA1001068	10386
	Y79AA1001077	F-Y79AA1001077	5469	R-Y79AA1001077	10387
	Y79AA1001078	F-Y79AA1001078	5470	R-Y79AA1001078	10388
	Y79AA1001105	F-Y79AA1001105	5471	R-Y79AA1001105	10389
35	Y79AA1001145	F-Y79AA1001145	5472	R-Y79AA1001145	10390
	Y79AA1001167	F-Y79AA1001167	5473	R-Y79AA1001167	10391
	Y79AA1001177	F-Y79AA1001177	5474	R-Y79AA1001177	10392
	Y79AA1001185	F-Y79AA1001185	5475	R-Y79AA1001185	10393
40	Y79AA1001211	F-Y79AA1001211	5476	R-Y79AA1001211	10394
	Y79AA1001216	F-Y79AA1001216	5477	R-Y79AA1001216	10395
	Y79AA1001228	F-Y79AA1001228	5478	R-Y79AA1001228	10396
	Y79AA1001233	F-Y79AA1001233	5479	R-Y79AA1001233	10397
45	Y79AA1001236	F-Y79AA1001236	5480	R-Y79AA1001236	10398
	Y79AA1001281	F-Y79AA1001281	5481	R-Y79AA1001281	10399
	Y79AA1001299	F-Y79AA1001299	5482	R-Y79AA1001299	10400
	Y79AA1001312	F-Y79AA1001312	5483	R-Y79AA1001312	10401
50	Y79AA1001323	F-Y79AA1001323	5484	R-Y79AA1001323	10402
	Y79AA1001384	F-Y79AA1001384	5485	R-Y79AA1001384	10403
	Y79AA1001391	F-Y79AA1001391	5486	R-Y79AA1001391	10404
	Y79AA1001394	F-Y79AA1001394	5487	R-Y79AA1001394	10405
55	Y79AA1001402	F-Y79AA1001402	5488	R-Y79AA1001402	10406
	Y79AA1001493	F-Y79AA1001493	5489	R-Y79AA1001493	10407

	Y79AA1001511	F-Y79AA1001511	5490	R-Y79AA1001511	10408
	Y79AA1001533	F-Y79AA1001533	5491	R-Y79AA1001533	10409
	Y79AA1001541	F-Y79AA1001541	5492	R-nnnnnnnnnnn	10410
5	Y79AA1001548	F-Y79AA1001548	5493	R-Y79AA1001548	10411
	Y79AA1001555	F-Y79AA1001555	5494	R-Y79AA1001555	10412
	Y79AA1001581	F-Y79AA1001581	5495		
	Y79AA1001585	F-Y79AA1001585	5496	R-Y79AA1001585	10413
10	Y79AA1001594	F-Y79AA1001594	5497	R-Y79AA1001594	10414
	Y79AA1001603	F-Y79AA1001603	5498	R-Y79AA1001603	10415
	Y79AA1001613	F-Y79AA1001613	5499	R-Y79AA1001613	10416
	Y79AA1001647	F-Y79AA1001647	5500	R-Y79AA1001647	10417
15	Y79AA1001665	F-Y79AA1001665	5501	R-Y79AA1001665	10418
	Y79AA1001679	F-Y79AA1001679	5502	R-Y79AA1001679	10419
	Y79AA1001692	F-Y79AA1001692	5503	R-nnnnnnnnnnn	10420
	Y79AÅ1001696	F-Y79AA1001696	5504	R-Y79AA1001696	10421
20	Y79AA1001705	F-Y79AA1001705	5505	R-Y79AA1001705	10422
	Y79AA1001711	F-Y79AA1001711	5506	R-Y79AA1001711	10423
	Y79AA1001781	F-Y79AA1001781	5507	R-Y79AA1001781	10424
	Y79AA1001805	F-Y79AA1001805	5508	R-nnnnnnnnnnn	10425
25	Y79AA1001827	F-Y79AA1001827	5509	R-Y79AA1001827	10426
23	Y79AA1001846	F-Y79AA1001846	5510	R-Y79AA1001846	10427
	Y79AA1001848	F-Y79AA1001848	5511	R-Y79AA1001848	10428
	Y79AA1001866	F-Y79AA1001866	5512	R-Y79AA1001866	10429
20	Y79AA1001874	F-Y79AA1001874	5513	R-Y79AA1001874	10430
30	Y79AA1001875	F-Y79AA1001875	5514	R-Y79AA1001875	10431
	Y79AA1001923	F-Y79AA1001923	5515	R-Y79AA1001923	10432
	Y79AA1001963	F-Y79AA1001963	5516		
	Y79AA1002027	F-Y79AA1002027	5517	R-Y79AA1002027	10433
35	Y79AA1002083	F-Y79AA1002083	5518	R-Y79AA1002083	10434
	Y79AA1002089	F-Y79AA1002089	5519	R-Y79AA1002089	10435
	Y79AA1002093	F-Y79AA1002093	5520	R-Y79AA1002093	10436
	Y79AA1002103	F-Y79AA1002103	5521	R-Y79AA1002103	10437
40	Y79AA1002115	F-Y79AA1002115	5522	R-Y79AA1002115	10438
	Y79AA1002125	F-Y79AA1002125	5523	R-Y79AA1002125	10439
	Y79AA1002139	F-Y79AA1002139	5524	R-Y79AA1002139	10440
	Y79AA1002204	F-Y79AA1002204	5525	R-Y79AA1002204	10441
45	Y79AA1002208	F-Y79AA1002208	5526	R-nnnnnnnnnn	10442
	Y79AA1002209	F-Y79AA1002209	5527	R-Y79AA1002209	10443
	Y79AA1002210	F-Y79AA1002210	5528	R-Y79AA1002210	10444
	Y79AA1002211	F-Y79AA1002211	5529	R-Y79AA1002211	10445
50	Y79AA1002220		5530	R-Y79AA1002220	10446
	Y79AA1002229	F-Y79AA1002229	5531	R-Y79AA1002229	10447
	Y79AA1002234		5532	R-Y79AA1002234	10448
	Y79AA1002246		5533	R-Y79AA1002246	10449
55	Y79AA1002258		5534	R-Y79AA1002258	10450
	Y79AA1002298	F-Y79AA1002298	5535	R-Y79AA1002298	10451

	Y79AA1002307	F-Y79AA1002307	5536	R-Y79AA1002307	10452
	Y79AA1002311	F-Y79AA1002311	5537	R-Y79AA1002311	10453
	Y79AA1002351	F-Y79AA1002351	5538	R-Y79AA1002351	10454
5	Y79AA1002361	F-Y79AA1002361	5539	R-Y79AA1002361	10455
	Y79AA1002399	F-Y79AA1002399	5540	R-Y79AA1002399	10456
	Y79AA1002407	F-Y79AA1002407	5541	R-Y79AA1002407	10457
10	Y79AA1002416	F-Y79AA1002416	5542	R-Y79AA1002416	10458
	Y79AA1002431	F-Y79AA1002431	5543	R-Y79AA1002431	10459
	Y79AA1002433	F-Y79AA1002433	5544	R-nnnnnnnnnnn	10460
	Y79AA1002472	F-Y79AA1002472	5545	R-Y79AA1002472	10461
15	Y79AA1002482	F-Y79AA1002482	5546	R-Y79AA1002482	10462
	Y79AA1002487	F-Y79AA1002487	5547	R-Y79AA1002487	10463

Table 2

5					OFO. TD
	name of	name of	SEQ ID		SEQ ID
	clone	5'-end	of 5'-end		of 3'-end
		sequence	sequence	sequence	sequence
10			_		·
	HEMBA1000497	F-HEMBA100049	7 16111	R-HEMBA100049	7 16165
	HEMBA1001750	F-HEMBA100175	0 16112	R-HEMBA100175	
45	HEMBA1003854	F-HEMBA100385	4 16113	R-HEMBA100385	4 16167
15	HEMBA1004193	F-НЕМВА100419	3 16114	R-HEMBA100419	
	HEMBA1004860	F-HEMBA100486	0 16115	R-HEMBA100486	
	HEMBA1005572	F-HEMBA100557	2 16116	R-HEMBA100557	
20	HEMBA1006038	F-HEMBA100603	8 16117	R-HEMBA100603	88 16171
	HEMBA1006092	F-HEMBA100609	2 16118	R-HEMBA100609	2 16172
	HEMBA1006406	F-HEMBA100640	6 16119	R-HEMBA100640	6 16173
	HEMBA1006650	F-HEMBA100665	0 16120	R-HEMBA100665	0 16174
25	HEMBA1006812	F-HEMBA100681	2 16121	R-HEMBA100681	
	HEMBB1000672	F-HEMBB100067	2 16122	R-HEMBB100067	
	HEMBB1001197	F-HEMBB100119	7 16123	R-HEMBB100119	
	HEMBB1001871	F-HEMBB100187	1 16124	R-HEMBB100187	
30	MAMMA1001252	F-MAMMA100125	2 16125	R-MAMMA100129	
	MAMMA1002094	F-MAMMA100209	4 16126	R-MAMMA100209	-
	NT2RM4000634	F-NT2RM400063	34 16127	R-NT2RM40006	
	NT2RM4000657	F-NT2RM400065	7 16128	R-NT2RM400069	
35	NT2RM4000783	F-NT2RM400078	33 16129	R-NT2RM400078	
	NT2RM4000857	F-NT2RM400085	7 16130	R-NT2RM40008	
	NT2RM4001178	F-NT2RM400117	78 16131	R-NT2RM40011	78 16185
40	NT2RM4002420	F-NT2RM400242	20 16132	R-NT2RM40024	20 16186
	NT2RP2000198	F-NT2RP200019	98 16133	R-NT2RP20001	98 16187
	NT2RP2000551	F-NT2RP20005	16134	R-NT2RP20005	51 16188
	NT2RP2000660	F-NT2RP20006	60 16135	R-NT2RP20006	60 16189

	NT2RP2001214	F-NT2RP2001214	16136	R-NT2RP2001214	16190
	NT2RP2001460	F-NT2RP2001460	16137	R-NT2RP2001460	16191
5	NT2RP2001756	F-NT2RP2001756	16138	R-NT2RP2001756	16192
3	NT2RP2002056	F-NT2RP2002056	16139	R-NT2RP2002056	16193
	NT2RP2002677	F-NT2RP2002677	16140	R-NT2RP2002677	16194
	NT2RP2002755	F-NT2RP2002755	16141	R-NT2RP2002755	16195
10	NT2RP2002843	F-NT2RP2002843	16142	R-NT2RP2002843	16196
	NT2RP2003101	F-NT2RP2003101	16143	R-NT2RP2003101	16197
	NT2RP2003799	F-NT2RP2003799	16144	R-NT2RP2003799	16198
	NT2RP2004095	F-NT2RP2004095	16145	R-NT2RP2004095	16199
15	NT2RP2004732	F-NT2RP2004732	16146	R-NT2RP2004732	16200
	NT2RP2004920	F-NT2RP2004920	16147	R-NT2RP2004920	16201
	NT2RP2005454	F-NT2RP2005454	16148	R-NT2RP2005454	16202
	NT2RP2005776	F-NT2RP2005776	16149	R-NT2RP2005776	16203
20	NT2RP2005806	F-NT2RP2005806	16150	R-NT2RP2005806	16204
	NT2RP2005882	F-NT2RP2005882	16151	R-NT2RP2005882	16205
	NT2RP3001282	F-NT2RP3001282	16152	R-NT2RP3001282	16206
25	NT2RP3001723	F-NT2RP3001723	16153	R-NT2RP3001723	16207
20	NT2RP3002099	F-NT2RP3002099	16154	R-NT2RP3002099	16208
	NT2RP3003155	F-NT2RP3003155	16155	R-NT2RP3003155	16209
	NT2RP3004028	F-NT2RP3004028	16156	R-NT2RP3004028	16210
30	OVARC1000008	F-0VARC1000008	16157	R-0VARC1000008	16211
	OVARC1000724	F-0VARC1000724	16158	R-0VARC1000724	16212
	OVARC1000751	F-0VARC1000751	16159	R-0VARC1000751	16213
	OVARC1001029	F-0VARC1001029	16160	R-OVARC1001029	16214
35	PLACE1000814	F-PLACE1000814	16161	R-PLACE1000814	16215
	PLACE1003030	F-PLACE1003030	16162	R-PLACE1003030	16216
	PLACE1005549	F-PLACE1005549	16163	R-PLACE1005549	16217
	PLACE1007218	F-PLACE1007218	16164	R-PLACE1007218	16218

[0019] Furthermore, the present invention relates to the use of the above primers, as described below.

- (4) A polynucleotide which can be synthesized with the primer set of (2) or (3).
- (5) A polynucleotide comprising a coding region in the polynucleotide of (4).
- (6) A substantially pure protein encoded by polynucleotide of (4).
- (7) A partial peptide of the protein of (6).

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[0020] In addition, the present invention comprises a polynucleotide described below and a protein encoded by the polynucleotide.

- (8) An isolated polynucleotide selected from the group consisting of
  - (a) a polynucleotide comprising a coding region of the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
    - (b) a polynucleotide comprising a nucleotide sequence encoding a protein comprising the amino acid sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
    - (c) a polynucleotide comprising a nucleotide sequence encoding a protein comprising an amino acid sequence

selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, in which one or more amino acids are substituted, deleted, inserted, and/or added, wherein said protein is functionally equivalent to the protein comprising said amino acid sequence selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351;

- (d) a polynucleotide that hybridizes with a polynucleotide comprising a nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351, and that comprises a nucleotide sequence encoding a protein functionally equivalent to the protein encoded by the nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351;
- (e) a polynucleotide comprising a nucleotide sequence encoding a partial amino acid sequence of a protein encoded by the polynucleotide of (a) to (d);
- (i) a polynucleotide comprising a nucleotide sequence with at least 70% identity to the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351.
- (9). A substantially pure protein encoded by the polynucleotide of (8).
- (10) An antibody against the protein or peptide of any one of (6), (7), and (9).
  - (11) A vector comprising the polynucleotide of (5) or (8).

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- (12) A transformant carrying the polynucleotide of (5) or (8), or the vector of (11).
- (13) A transformant expressively carrying the polynucleotide of (5) or (8), or the vector of (11).
- (14) A method for producing the protein or peptide of any one of (6), (7), and (9), comprising culturing the transformant of (13) and recovering the expression product.
- (15) An oligonucleotide comprising the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351 or the nucleotide sequence complementary to the complementary strand thereof, wherein said oligonucleotide comprises 15 nucleotides or more.
- (16) Use of the oligonucleotide of (15) as a primer for synthesizing a polynucleotide.
- (17) Use of the oligonucleotide of (15) as a probe for detecting a gene.
- (18) An antisense polynucleotide against the polynucleotide of (8), or the portion thereof.
- (19) A method for synthesizing a polynucleotide, the method comprising:
  - a) synthesizing a complementary strand using a cDNA library as a template, and using the primer set of (2) or (3), or the primer of (16); and
  - b) recovering the synthesized product.
- (20) The method of (19), wherein the cDNA library is obtainable by oligo-capping method.
- (21) The method of (19), wherein the complementary strand is obtainable by PCR.
- (22) A method for detecting the polynucleotide of (8), the method comprising:
  - a) incubating a target polynucleotide with the oligonucleotide of (15) under the conditions where hybridization occurs, and
  - b) detecting the hybridization of the target polynucleotide with the oligonucleotide of (15).
  - (23) A database of polynucleotides and/or proteins, the database comprising information on at least one sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351 and/or the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, or a medium on which the database is stored.
- 45 [0021] Any patents, patent applications, and publications cited herein are incorporated by reference.

## BRIEF DESCRIPTION OF THE DRAWINGS

- [0022] Figure 1 shows the restriction maps of vectors pME18SFL3 and pUC19FL3.
- Figure 2 shows the reproducibility of gene expression analysis. The respective intensities of gene expression observed in independent set of experiments are plotted in the vertical axis as well as in the horizontal axis.
  - [0024] Figure 3 shows the detection limit in gene expression analysis. The intensity of expression is shown in the vertical axis and the concentration ( $\mu$ g/ml) of probe used is shown in the horizontal axis.

### 55 DETAILED DESCRIPTION OF THE INVENTION

[0025] Herein, "polynucleotide" is defined as a molecule in which multiple nucleotides are polymerized. There are no limitations in the number of the polymerized nucleotides. In case that the polymer contains relatively low number

of nucleotides, it is also described as an "oligonucleotide". The polynucleotide or the oligonucleotide of the present invention can be a natural or chemically synthesized product. Alternatively, it can be synthesized using a template DNA by an enzymatic reaction such as PCR.

[0026] All the cDNA provided by the invention are full-length cDNA. Herein, a "full-length cDNA" is defined as a cDNA which contains both ATG codon (the translation start site) and the stop codon. Accordingly, the untranslated regions, which are originally found in the upstream or downstream of the protein coding region in natural mRNA, may or may not be contained.

[0027] An "isolated polynucleotide" is a polynucleotide the structure of which is not identical to that of any naturally occurring nucleic acid or to that of any fragment of a naturally occurring genomic nucleic acid spanning more than three separate genes. The term therefore covers, for example,

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- (a) a DNA which has the sequence of part of a naturally occurring genomic DNA molecule but is not flanked by both of the coding sequences that flank that part of the molecule in the genome of the organism in which it naturally occurs:
- (b) a nucleic acid incorporated into a vector or into the genomic DNA of a prokaryote or eukaryote in a manner such that the resulting molecule is not identical to any naturally occurring vector or genomic DNA;
- (c) a separate molecule such as a cDNA, a genomic fragment, a fragment produced by polymerase chain reaction (PCR), or a restriction fragment; and
- (d) a recombinant nucleotide sequence that is part of a hybrid gene, i.e., a gene encoding a fusion protein. Specifically excluded from this definition are nucleic acids present in mixtures of different (i) DNA molecules, (ii) transfected cells, or (iii) cell clones: e.g., as these occur in a DNA library such as a cDNA or genomic DNA library.

[0028] The term "substantially pure" as used herein in reference to a given polypeptide means that the protein or polypeptide is substantially free from other biological macromolecules. The substantially pure protein or polypeptide is at least 75% (e.g., at least 80, 85, 95, or 99%) pure by dry weight. Purity can be measured by any appropriate standard method, for example, by column chromatography, polyacrylamide gel electrophoresis, or HPLC analysis.

[0029] All the clones (5602 clones) of the present invention are novel and encode the full-length proteins. All the clones were prepared by oligo capping method, which can achieve cDNA cloning with high fullness ratio. The cDNA clones were selected by using ATGprI score as an index of the fullness ratio at the 5'-end, based on the sequence features of the 5'-end sequences. Selection was further carried out by searching GenBank database for EST sequences homologous to 5'-end sequence of each clone by BLAST [S.F. Altschul, W. Gish, W. Miller, E.W. Myers & D.L. Lipman J. Mol. Biol., 215:403-410 (1990); W. Gish, & D.J. States, Nature Genet., 3:266-272 (1993)] and by considering the number of matching (identical) EST sequences or the number of continuous amino acids in the 5'-end sequence initiated from the initiation codon.

[0030] Moreover, the clones were turn out to be not identical to any of the known human mRNA (namely novel) by homology search using the 5'-end sequence.

[0031] The primers of the present invention, which are used for synthesizing full-length cDNA, are selected from the group comprising SEQ ID NO: 1-5547 (5'-primer), or SEQ ID NO: 5548-10463 (3'-primer). Further, the primers of the present invention, which are used for synthesizing full-length cDNA, are selected from SEQ ID NO: 16111-16164 (5'-primer), or SEQ ID NO: 16165-16218 (3'-primer). Some of the nucleotides include a known EST as its part. However, the primers of the present invention are novel in terms that the primers enable to synthesize full-length cDNA. Because the known ESTs lack important information on what part of cDNA the ESTs correspond to, it is impossible to design primers on the basis of the ESTs.

[0032] All the full-length cDNA of the present invention can be synthesized using a primer set comprising the nucleotide sequences selected from both the 5'-and 3'-end sequences, or a set comprising a primer based on the 5'-end sequence and an oligo-dT primer, by a method such as PCR (Current protocols in Molecular Biology (1987) Ausubel et al. edit, John Wiley & Sons, Section 6.1-6.4).

[0033] Specifically, PCR can be performed using an oligonucleotide that has 15 nucleotides longer, and specifically hybridizes with the complementary strand of the polynucleotide that contains the nucleotide sequence selected from the 5'-end sequences shown in Table 1 and 2 (SEQ ID NO: 1-5547, or SEQ ID NO: 16111-16164), and an oligo-dT primer as a 5'-, and 3'-primer, respectively. The length of the primers is usually 15-100 bp, and favorably between 15-35 bp. In case of LA PCR, which is described below, the primer length of 25-35 bp may provide a good result.

[0034] A method to design a primer that enables a specific amplification based on the given nucleotide sequence is known to those skilled in the art (Current Protocols in Molecular Biology, Ausubel et al. edit, (1987) John Wiley & Sons, Section 6.1-6.4). In designing a primer based on the 5'-end sequence, the primer is designed so as that, in principle, the amplification products will include the translation start site. Accordingly, in case that a given 5'-end nucleotide sequence is the 5'- untranslated region (5'UTR), any part of the sequence can be used as a 5'-primer as far as the specificity toward the target cDNA is insured. The translation start site can be predicted using a known method such

as the ATGpr as described below.

[0035] When synthesizing a polynucleotide, the target nucleotide sequence to be amplified can extend to several thousand bp in some cDNA. However, it is possible to amplify such a long nucleotides by using such as LA PCR (Long and Accurate PCR). It is advantageous to use LA PCR when synthesizing long DNA. In LA PCR, in which a special DNA polymerase having 3' ->5' exonuclease activity is used, misincorporated nucleotides can be removed. Accordingly, accurate synthesis of the complementary strand can be achieved even with a long nucleotide sequence. By using LA PCR, it is reported that amplification of a nucleotide with 20 kb longer can be achieved under desirable condition (Takeshi Hayashi (1996) Jikken-Igaku Bessatsu, "Advanced Technologies in PCR" Youdo-sha).

[0036] A template DNA for synthesizing the cDNA of the present invention can be obtained by using cDNA libraries that are prepared by various methods. The full-length cDNA clones obtained here are those with high fullness ratio, which were obtained using a combination of (1) a method to prepare a full-length-enriched cDNA library using the oligo-capping method, and (2) an estimation system for fullness using the 5'-end sequence (selection based on the estimation by the ATGpr after removing clones that are not-full-length compared to the ESTs). However, it is possible to easily obtain a full-length cDNA by using the primers that are provided by the present invention, not by the above described specialized method.

The problem with the cDNA libraries prepared by the known methods or commercially available is that mRNA contained in the libraries has very low fullness ratio. Thus, it is difficult to screen full-length cDNA clone directly from the library using ordinary cloning methods. The present invention has revealed a primer that is capable of synthesizing a full-length cDNA. If provided with primers, it is possible to synthesize a target full-length cDNA by using enzymatic reactions such as PCR. In particular, a full-length-enriched cDNA library, synthesized by methods such as oligo-capping, is desirable to synthesize a full-length cDNA with more reliability.

[0037] The 5'-end sequence of the full-length cDNA clones of the invention can be used to isolate the regulatory element of transcription including the promoter on the genome. By the spring of the year 2000, a rough draft of the human genome (analysis of human genomic sequence with lower accuracy), which covers 90% of the genome, is planned to be accomplished, and by the year 2003, analysis of the entire human genomic sequence is going to be finished. However, it is hard to analyze with software the transcription start sites on the human genome, in which long introns exist. By contrast, it is easy to specify the transcription start site on the genomic sequence using the 5'-end sequence of the full-length cDNA clone, thus it is easy to obtain the genomic region involved in transcription regulation, which includes the promoter that is contained in the upstream of the transcription start site.

[0038] The full-length cDNAs cloned in the present invention are classified into 13 groups, based on the data such as ATGpr1 score, by which the fullness ratio can be evaluated. Specifically, the 13 groups consist of; the below-mentioned groups (1)-(3), containing 3690 clones (Table 9), and the group (12), containing 3 clones, wherein ATGpr1 (score defined in the ATGpr program) is higher than 0.3; and the below-mentioned groups (4)-(11), containing 1857 clones (Table 10), and the group (13), containing 52 clones, wherein, although ATGpr1 is 0.3 or less, the clones are judged to be full-length from various viewpoints. Names of the clones belonging to the groups (1)-(13) are as indicated in Examples or below.

### (1) 1516 clones

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Among the 3690 clones that have the maximal ATGpr1 score higher than 0.3, 1516 clones are novel full-length clones, in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

### (2)377 clones

Among the 3690, 377 clones are novel full-length clones, in which the number of human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

## (3) 1797 clones

Among the 3690, 1797 clones are novel full-length clones, in which the number of human EST having identical sequence at the 5'-end is not more than 20 (except the clones described above).

### (4) 453 clones

Among the 1857 clones in which the maximal ATGpr1 score is 0.3 or less, the following 453 clones are estimated to be novel full-length clones since the clones have the maximal score 0.3 or more in the ATGpr2, and at least either of the sequences of their 5'- and 3'-ends, or both are not identical to those of any human EST. The ATGpr2 score is determined by using the ATGpr program with neglecting the information of the frequency of the six nucleotides contained within the sequence between the ATG codon and the stop codon (the maximal length is 300 nucleotides from the ATG codon) (Salamo A.A., Nishikawa T., and Swindells M.B. (1998) Bioinformatics, 14: 384-390; http://www.hri.co.jp/atgpr/). The ATGpr program for calculating the ATGpr2 score is described as the

ATGpr2 program in the followings.

#### (5) 24 clones

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Among the 1857 clones, 24 clones are estimated to be full-length since their maximal ATGpr2 scores are higher than 0.3, and also novel, though they have low scores in ATGpr1 program, in which the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

## (6) 65 clones

Among the 1857 clones, 65 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

### (7) 32 clones

Among the 1857 clones, 32 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

## (8) 36 clones

Among the 1857 clones, 36 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

### (9) 81 clones

Among the 1857 clones, 81 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if the number of the human EST having identical sequence at the 5'-end is not more than 20 (other than the clones in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST).

### (10) 938 clones

Among the 1857 clones, 938 clones are estimated to be full-length according to the fullness ratio shown in Table 4, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 5'-end is not identical to those of any human EST.

### (11) 228 clones

Among the 1857 clones, 228 clones are estimated to be full-length according to the fullness ratio shown in Table 7, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 3'-end is not identical to those of any human EST.

### (12) 3 clones

Three clones, HEMBA1006812, HEMBB1001871, and NT2RP3001282, whose maximal ATGpr1 values are higher than 0.3, are full-length and novel clones whose 5'-end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 100 amino acids or more.

### (13) 52 clones

The following 52 clones, which have maximal ATGpr1 values of 0.3 or less, are full-length with the fullness ratios shown in Table 4 although the fullness ratios are low:

HEMBA1000497,	HEMBA1001750,	HEMBA1003854,	HEMBA1004193,	HEMBA1004860,	HEMBA1005572,
HEMBA1006038,	HEMBA1006092,	HEMBA1006406,	HEMBA1006650,	HEMBB1000672,	HEMBB1001197,
MAMMA1001252,	MAMMA1002094,	NT2RM4000634,	NT2RM4000657,	NT2RM4000783,	NT2RM4000857,
NT2RM4001178,	NT2RM4002420,	NT2RP2000198,	NT2RP2000551,	NT2RP2000660,	NT2RP2001214,
NT2RP2001460,	NT2RP2001756,	NT2RP2002056,	NT2RP2002677,	NT2RP2002755,	NT2RP2002843,
NT2RP2003101,	NT2RP2003799,	NT2RP2004095,	NT2RP2004732,	NT2RP2004920,	NT2RP2005454,
NT2RP2005776,	NT2RP2005806,	NT2RP2005882,	NT2RP3001723,	NT2RP3002099,	NT2RP3003155,
NT2RP3004028.	OVARC1000008.	OVARC1000724,	OVARC1000751,	OVARC1001029,	PLACE1000814,

PLACE1003030, PLACE1005549, PLACE1007218, NT2RP4002298.

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Moreover, the clones are novel clones whose 5' -end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 50 amino acids or more. Among them, the following 20 clones is predicted to contain a coding region with 100 amino acids or more and should encode proteins:

HEMBA100497, HEMBA1003854, HEMBA1004193, NT2RM4000657, NT2RM4001178, NT2RP2001756, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2004095, NT2RP2004920, NT2RP3002099, NT2RP3003155, OVARC1000724, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218.

[0039] The protein encoded by the polynucleotide of the invention can be prepared as a recombinant protein or as a natural protein. For example, the recombinant protein can be prepared by inserting the polynucleotide encoding the protein of the invention into a vector, introducing the vector into an appropriate host cell and purifying the protein expressed within the transformed host cell, as described below. In contrast, the natural protein can be prepared, for example, by utilizing an affinity column to which an antibody against the protein of the invention (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 16.1-16.19) is attached. The antibody used for affinity purification may be either a polyclonal antibody, or a monoclonal antibody. Alternatively, in vitro translation (See, for example, "On the fidelity of mRNA translation in the nuclease-treated rabbit reticulocyte lysate system." Dasso M.C., and Jackson R.J. (1989) Nucleic Acids Res. 17: 3129-3144) may be used for preparing the protein of the invention.

[0040] Proteins functionally equivalent to the proteins of the present invention can be prepared based on the activities, which were clarified in the above-mentioned manner, of the proteins of the present invention. Using the biological activity possessed by the protein of the invention as an index, it is possible to verify whether or not a particular protein is functionally equivalent to the protein of the invention by examining whether or not the protein has said activity.

[0041] Proteins functionally equivalent to the proteins of the present invention can be prepared by those skilled in the art, for example, by using a method for introducing mutations into an amino acid sequence of a protein (for example, site-directed mutagenesis (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 8.1-8.5). Besides, such proteins can be generated by spontaneous mutations. The present invention comprises the proteins having one or more amino acids substitutions, deletions, insertions and/or additions in the amino acid sequences of the proteins of the present invention (Tables 350 and 351), as far as the proteins have the equivalent functions to those of the proteins identified in the present Examples described later.

[0042] There are no limitations in the number and sites of amino acid mutations, as far as the proteins maintain the functions thereof. The number of mutations is typically 30% or less, or 20% or less, or 10% or less, preferably within 5% or less, or 3% or less of the total amino acids, more preferably within 2% or less or 1 % or less of the total amino acids. From the viewpoint of maintaining the protein function, it is preferable that a substituted amino has a similar property to that of the original amino acid. For example, Ala, Val, Leu, Ile, Pro, Met, Phe and Trp are assumed to have similar properties to one another because they are all classified into a group of non-polar amino acids. Similarly, substitution can be performed among non-charged amino acid such as Gly, Ser, Thr, Cys, Tyr, Asn, and Gln, acidic amino acids such as Asp and Glu, and basic amino acids such as Lys, Arg, and His.

[0043] In addition, proteins functionally equivalent to the proteins of the present invention can be isolated by using techniques of hybridization or gene amplification known to those skilled in the art. Specifically, using the hybridization technique (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)), those skilled in the art can usually isolate a DNA highly homologous to the DNA encoding the protein identified in the present Example based on the identified nucleotide sequence (Tables 350 and 351) or a portion thereof and obtain the functionally equivalent protein from the isolated DNA. The present invention include proteins encoded by the DNAs hybridizing with the DNAs encoding the proteins identified in the present Example, as far as the proteins are functionally equivalent to the proteins identified in the present Example. Organisms from which the functionally equivalent proteins are isolated are illustrated by vertebrates such as human, mouse, rat, rabbit, pig and bovine, but are not limited to these animals.

[0044] Washing conditions of hybridization for the isolation of DNAs encoding the functionally equivalent proteins are usually "1 × SSC, 0.1% SDS, 37°C"; more stringent conditions are "0.5 × SSC, 0.1% SDS, 42°C"; and still more stringent conditions are "0.1 × SSC, 0.1% SDS, 65°C". Alternatively, the following conditions can be given as hybridization conditions of the present invention. Namely, conditions in which the hybridization is done at "6 × SSC, 40% Formamide, 25°C", and the washing at "1 × SSC, 55°C" can be given. More preferable conditions are those in which the hybridization is done at "6 × SSC, 40% Formamide, 37°C", and the washing at "0.2 × SSC, 55°C". Even more preferable are those in which the hybridization is done at "6 × SSC, 50% Formamide, 37°C", and the washing at "0.1 × SSC, 62°C". The more stringent the conditions of hybridization are, the more frequently the DNAs highly homologous to the probe sequence are isolated. Therefore, it is preferable to conduct hybridization under stringent conditions. Examples of stringent conditions in the present invention are, washing conditions of "0.5 × SSC, 0.1% SDS, 42°C", or alternatively, hybridization conditions of "6 × SSC, 40% Formamide, 37°C", and the washing at "0.2 × SSC, 55°C". However, the above-mentioned combinations of SSC, SDS and temperature conditions are indicated just as examples.

Those skilled in the art can select the hybridization conditions with similar stringency to those mentioned above by properly combining the above-mentioned or other factors (for example, probe concentration, probe length and duration of hybridization reaction) that determines the stringency of hybridization.

[0045] The amino acid sequences of proteins isolated by using the hybridization techniques usually exhibit high homology to those of the proteins of the present invention, which are shown in Tables 350 and 351. The present invention encompasses a polynucleotide comprising a nucleotide sequence that has a high identity to the nucleotide sequence of claim 8 (a).

Furthermore, the present invention encompasses a peptide, or protein comprising an amino acid sequence that has a high identity to the amino acid sequence encoded by the polynucleotide of claim 8 (b). The term "high identity" indicates sequence identity of at least 40% or more;

preferably 60% or more; and more preferably 70% or more. Alternatively, more preferable is identity of 90% or more, or 93% or more, or 95% or more, furthermore, 97% or more, or 99% or more. The identity can be determined by using the BLAST search algorithm.

[0046] With the gene amplification technique (PCR) (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)) using primers designed based on the nucleotide sequence (Tables 350 and 351) or a portion thereof identified in the present Example, it is possible to isolate a DNA fragment highly homologous to the polynucleotide sequence or a portion thereof and to obtain functionally equivalent protein to a particular protein identified in the present Example based on the isolated DNA fragment.

[0047] The "percent identity" of two amino acid sequences or of two nucleic acids is determined using the algorithm of Karlin and Altschul (Proc. Natl. Acad. Sei. USA 87:2264-2268, 1990), modified as in Karlin and Altschul (Proc. Natl. Acad. Sei. USA 90:5873-5877, 1993). Such an algorithm is incorporated into the BLASTN and BLASTX programs of Altschul et al. (J. Mol. Biol.215:403-410, 1990). BLAST nucleotide searches are performed with the BLASTN program, score = 100, wordlength = 12. BLAST protein searches are performed with the BLASTX program, score = 50, wordlength = 3. When gaps exist between two sequences, Gapped BLAST is utilized as described in Altschul et al. (Nucleic Acids Res.25:3389-3402,1997). When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (e.g., BLASTX and BLASTN) are used. See http://www.ncbi.nlm.nih.gov.

[0048] The present invention also includes a partial peptide of the proteins of the invention. The partial peptide comprises a protein generated as a result that a signal peptide has been removed from a secretory protein. If the protein of the present invention has an activity as a receptor or a ligand, the partial peptide may function as a competitive inhibitor of the protein and may bind to the receptor (or ligand). In addition, the present invention comprises an antigen peptide for raising antibodies. For the peptides to be specific for the protein of the invention, the peptides comprise at least 7 amino acids, preferably 8 amino acids or more, more preferably 9 amino acids or more, and even more preferably 10 amino acids or more. The peptide can be used for preparing antibodies against the protein of the invention, or competitive inhibitors of them, and also screening for a receptor that binds to the protein of the invention. The partial peptides of the invention can be produced, for example, by genetic engineering methods, known methods for synthesizing peptides, or digesting the protein of the invention with an appropriate peptidase.

[0049] The present invention also relates to a vector into which the DNA of the invention is inserted. The vector of the invention is not limited as long as it contains the inserted DNA stably. For example, if *E. coli* is used as a host, vectors such as pBluescript vector (Stratagene) are preferable as a cloning vector. To produce the protein of the invention, expression vectors are especially useful. Any expression vector can be used as far as it is capable of expressing the protein *in vitro*, in *E. coli*, in cultured cells, or *in vivo*. For example, pBEST vector (Promega) is preferable for *in vitro* expression, pET vector (Invitrogen) for *E. coli*, pME18S-FL3 vector (GenBank Accession No. AB009864) for cultured cells, and pME18S vector (Mol. Cell. Biol. (1988) 8: 466-472) for *in vivo* expression. To insert the DNA of the invention, ligation utilizing restriction sites can be performed according to the standard method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

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[0050] The present invention also relates to a transformant carrying the vector of the invention. Any cell can be used as a host into which the vector of the invention is inserted, and various kinds of host cells can be used depending on the purposes. For strong expression of the protein in eukaryotic cells, COS cells or CHO cells can be used, for example. [0051] Introduction of the vector into host cells can be performed, for example, by calcium phosphate precipitation method, electroporation method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 9.1-9.9), lipofectamine method (GIBCO-BRL), or microinjection method, etc.

[0052] The primer of the present invention can be used for synthesizing full-length cDNA, and also for the detection and/or diagnosis of the abnormality of the protein of the invention encoded by the full-length cDNA. For example, by utilizing polymerase chain reaction (genomic DNA-PCR, or RT-PCR) using the primer of the invention, DNA encoding the protein of the invention can be amplified. It is also possible to obtain the regulatory region of expression in the 5'-upstream by using PCR or hybridization since the transcription start site within the genomic sequence can be easily specified based on the 5'-end sequence of the full-length cDNA. The obtained genomic region can be used for detection and/or diagnosis of the abnormality of the sequence by RFLP analysis, SSCP, or direct sequencing.

[0053] Furthermore, the "polynucleotide having a length of at least 15 nucleotides, comprising a nucleotide sequence that is complementary to a polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs in Tables 350 and 351, or its complementary strand" includes an antisense polynucleotide for suppressing the expression of the protein of the invention. To exert the antisense effect, the antisense polynucleotide has a length of at least 15 bp or more, for example, 50 bp or more, preferably 100 bp or more, and more preferably 500 bp or more, and has a length of usually 3000 bp or less and preferably 2000 bp or less. The antisense DNA can be used in the gene therapy of the diseases that are caused by the abnormality of the protein of the invention (abnormal function or abnormal expression). Said antisense DNA can be prepared, for example, by the phosphorothioate method ("Physicochemical properties of phosphorothioate oligodeoxynucleotides." Stein (1988) Nucleic Acids Res. 16: 3209-3221) based on the nucleotide sequence of the DNA encoding the protein (for example, the DNA set forth in any one of SEQ ID NOs in Tables 350 and 351).

[0054] The polynucleotide or antisense DNA of the present invention can be used in gene therapy, for example, by administrating it into a patient by the in vivo or ex vivo method with virus vectors such as retrovirus vectors, adenovirus vectors, and adeno-associated virus vectors, or non-virus vectors such as liposome.

[0055] The present invention also relates to antibodies that bind to the protein of the invention.

There are no limitations in the form of the antibodies of the invention. They include polyclonal antibodies, monoclonal antibodies, or their portions that can bind to the protein of the invention. They also include antibodies of all classes. Furthermore, special antibodies such as humanized antibodies are also included.

[0056] The polyclonal antibody of the invention can be obtained according to the standard method by synthesizing an oligopeptide corresponding to the amino acid sequence and immunizing rabbits with the peptides (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.12-11.13). The monoclonal antibody of the invention can be obtained according to the standard method by purifying the protein expressed in *E. coli*, immunizing mice with the protein, and producing a hybridoma cell by fusing the spleen cells and myeloma cells (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

[0057] The antibody binding to the protein of the present invention can be used for purification of the protein of the invention, and also for detection and/or diagnosis of the abnormalities of the expression and structure of the protein. Specifically, proteins can be extracted, for example, from tissues, blood, or cells, and the protein of the invention is detected by Western blotting, immunoprecipitation, or ELISA, etc. for the above purpose.

[0058] Furthermore, the antibody binding to the protein of the present invention can be utilized for treating the diseases that associates with the protein of the invention. If the antibodies are used for treating patients, human antibodies or humanized antibodies are preferable in terms of their low antigenicity. The human antibodies can be prepared by immunizing a mouse whose immune system is replaced with that of human ("Functional transplant of megabase human immunoglobulin loci recapitulates human antibody response in mice" Mendez M.J. et al. (1997) Nat. Genet. 15: 146-156). The humanized antibodies can be prepared by recombination of the hypervariable region of a monoclonal antibody (Methods in Enzymology (1991) 203: 99-121).

[0059] The cDNA of the present invention encodes the amino acid sequence of a protein which is predicted to have the function(s) described below based on the homology search of the GenBank and SwissProt. Specifically, for instance, as shown in EXAMPLES, searching a known gene or protein that is homologous to the partial sequence of the full-length cDNA of the invention (5602 clone) and referring the function of the gene and of the protein encoded by the gene make it possible to predict the function of the protein encoded by the cDNA of the invention. In this way, each of 1437 clones out of the 5602 full-length cDNA clones of the invention was predicted to encode a protein that was classified into one or more of the following categories.

Secretory or membrane protein (261 clones)

Glycoprotein-associated protein (113 clones)

Signal transduction-associated protein (148 clones)

Transcription-associated protein (233 clones)

Disease-associated protein (437 clones)

Enzyme or metabolism-associated protein (301 clones)

Cell division- or cell proliferation-associated protein (74 clones)

Cytoskeleton-associated protein (92 clones)

RNA synthesis-associated protein (280 clones)

Nuclear protein (352 clones)

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Protein synthesis- or transport-associated protein (112 clones)

Cellular defense-associated protein (23 clones)

Development- or growth-associated protein (23 clones)

[0060] It is also possible to predict the protein function by looking into the amino acid sequence for the motifs such

as the signal sequence, transmembrane region, nuclear translocation signal, glycosylation signal, phosphorylation site, Zinc finger motif, and SH3 domain. The programs, PSORT (Nakai K., and Kanehisa M. (1992) Genomics 14: 897-911), SOSUI (Hirokawa T. et al. (1998) Bioinformatics 14: 378-379) (Mitsui Information Developing Inc.), and MEMSAT (Jones D.T., Taylor W.R., and Thornton J.M. (1994) Biochemistry 33: 3038-3049) can be used to predict the existence of the signal sequence or transmembrane region. Alternatively, a partial amino acid sequence of the protein is fused with another protein such as GFP, the fusion protein is transfected into cultured cells, and the localization is analyzed to predict the function of the original protein.

[0061] Based on the determined nucleotide sequences of the full-length cDNAs obtained in the present invention, it is possible to predict more detailed functions of the proteins encoded by the cDNA clones, for example, by searching the databases such as GenBank, Swiss-Prot and UniGene for homologies of the cDNAs; or by searching the amino acid sequences deduced from the full-length cDNAs for signal sequences by using software programs such as PSORT, for transmembrane regions by using software programs such as SOSUI or for motifs by using software programs such as Pfam (http://www.sanger.ac.uk/Software/Pfam/index.shtm1) and PROSITE (http://www.expasy.ch/prosite/). As a matter of course, the functions are often predictable by using partial sequence information (preferably 300 nucleotides or more) instead of the full-length nucleotide sequences. However, the result of the prediction by using partial nucleotide sequence does not always agree with the result obtained by using full-length nucleotide sequence, and thus, it is needless to say that the prediction of function is preferably performed based on the full-length nucleotide sequences. GenBank, Swiss-Prot and UniGene databases were searched for homologies of the full-length nucleotide sequences of the 4997 clones (see Example 18). The amino acid sequences deduced from the full-length nucleotide sequences were searched for functional domains by PSORT, SOSUI and Pfam. Prediction of functions of proteins encoded by the clones and the categorization thereof were performed based on these results obtained.

The following 798 clones were categorized into secretory and/or membrane proteins.

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HEMBA1000356, HEMBA1000518, HEMBA1000531, HEMBA1000637,
                                                          HEMBA1000719, HEMBA1000817,
                                                          HEMBA1001052, HEMBA1001071,
HEMBA1000822.
              HEMBA1000852,
                            HEMBA1000870, HEMBA1000991,
                                                          HEMBA1001446, HEMBA1001515,
HEMBA1001085.
              HEMBA1001286,
                             HEMBA1001351, HEMBA1001407,
                             HEMBA1001661, HEMBA1001734,
                                                          HEMBA1001746, HEMBA1001866,
HEMBA1001557,
              HEMBA1001569,
                             HEMBA1002166, HEMBA1002417,
                                                          HEMBA1002462, HEMBA1002475,
HEMBA1002125,
              HEMBA1002150,
                                                          HEMBA1002661, HEMBA1002780,
HEMBA1002477.
              HEMBA1002486,
                             HEMBA1002609, HEMBA1002659,
                             HEMBA1002921, HEMBA1003071,
                                                                        HEMBA1003079,
                                                          HEMBA1003077,
HEMBA1002818,
              HEMBA1002876,
                             HEMBA1003281, HEMBA1003286,
                                                          HEMBA1003538,
                                                                        HEMBA1003711,
HEMBA1003086,
              HEMBA1003096,
                             HEMBA1004055, HEMBA1004143,
                                                          HEMBA1004146,
                                                                         HEMBA1004207.
HEMBA1003742,
              HEMBA1003803,
                                           HEMBA1004637,
                                                          HEMBA1004752,
                                                                         HEMBA1004756,
HEMBA1004341,
              HEMBA1004461,
                             HEMBA1004577,
              HEMBA1004889.
                             HEMBA1004923.
                                           HEMBA1004930,
                                                          HEMBA1005029,
                                                                         HEMBA1005035,
HEMBA1004850,
                                                                         HEMBA1005616,
HEMBA1005050.
              HEMBA1005552,
                             HEMBA1005576,
                                            HEMBA1005581,
                                                          HEMBA1005588,
HEMBA1005699,
              HEMBA1005991,
                             HEMBA1006036,
                                           HEMBA1006038,
                                                          HEMBA1006067.
                                                                         HEMBA1006173,
HEMBA1006198,
              HEMBA1006293,
                             HEMBA1006310,
                                            HEMBA1006492,
                                                          HEMBA1006502,
                                                                         HEMBA1006583,
              HEMBA1006758,
                             HEMBA1006789,
                                            HEMBA1006921.
                                                          HEMBA1006926,
                                                                         HEMBA1006976,
HEMBA1006659,
HEMBA1007203,
              HEMBA1007301,
                             HEMBB1000037.
                                            HEMBB1000050,
                                                          HEMBB1000054,
                                                                         HEMBB1000175,
HEMBB1000317,
              HEMBB1000556,
                             HEMBB1000593,
                                            HEMBB1000631,
                                                          HEMBB1000763,
                                                                         HEMBB1000827.
HEMBB1000915,
              HEMBB1000975,
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    [0062] The following 142 clones were categorized into glycoprotein-associated proteins.
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    HEMBA1003679, HEMBA1003866, HEMBA1005576, HEMBA1005581, HEMBA1005699,
    HEMBA1006976, HEMBA1007301, HEMBB1000317, HEMBB1000915, HEMBB1001871,
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    HEMBA1005737, HEMBA1006130, HEMBA1006708, HEMBB1000083, HEMBB1000266, HEMBB1000632,
    HEMBB1000781, HEMBB1000831, HEMBB1002193, MAMMA1000173, MAMMA1001038,
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     [0066] The following 427 clones presumably belong to enzymes and/or metabolism-associated proteins.
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                                                                                  OVARC1001180,
                    OVARC1000326,
                                   OVARC1000556,
    OVARC1000241.
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    OVARC1001232,
                    OVARC1001271,
                                   OVARC1001306,
                                                   OVARC1001436,
                                                                   OVARC1002112,
                                                                                   PLACE1001118,
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                    PLACE4000326,
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     THYRO1000585,
                    THYRO1001100,
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                    Y79AA1000589,
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     Y79AA1001963, Y79AA1002431, Y79AA1002472, Y79AA1002482,
     [0069] The following 296 clones presumably belong to DNA- and/or RNA-binding proteins. HEMBA1000158,
                                   HEMBA1000591,
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                                                                  HEMBA1001819,
                                                                                  HEMBA1001847.
    HEMBA1001405,
                    HEMBA1001510.
                                   HEMBA1001804.
                                                                                  HEMBA1003568,
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                                                   HEMBA1003408,
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35
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                                   HEMBA1004354,
                                                   HEMBA1004389,
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                                                   HEMBB1001482,
                                                                  HEMBB1001736, HEMBB1001749,
     HEMBB1000264,
                    HEMBB1000789.
                                                                  MAMMA1000731, MAMMA1001105,
     HEMBB1001839, HEMBB1002217, MAMMA1000183, MAMMA1000284,
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     MAMMA1001222, MAMMA1001260, MAMMA1001743, MAMMA1001837,
     MAMMA1002869, MAMMA1002937, MAMMA1003011, NT2RM1000086, NT2RM1000539, NT2RM1000555,
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45
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                                                   OVARC1001232,
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                                   PLACE1003605.
                                                   PLACE1003704.
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     HEMBA1004596, OVARC1000148, PLACE1003334, THYRO1001661,
    [0070] The following 66 clones presumably belong to the category of RNA synthesis-associated proteins.
    HEMBA1000591, HEMBA1001579, HEMBA1003179, HEMBA1003591, HEMBA1006278, HEMBB1000226,
                                                  NT2RM2001989.
     NT2RM1000187. NT2RM1000852.
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                                                                  NT2RM2002100, NT2RM4000191,
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    NT2RM4001178.
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                    NT2RP3000397,
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                                                                                  PLACE1001739.
     PLACE1003704,
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                                                   PLACE1004814,
                                                                  PLACE1004902,
                                                                                  PLACE1005373,
    PLACE1005646.
                    PLACE1005876,
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                                                   PLACE1006626,
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                                                                                  PLACE1006917.
    PLACE1009476, PLACE1009925, PLACE1010194, PLACE1011114, THYRO1000121, Y79AA1001963,
     [0071] The following 184 clones presumably belong to protein synthesis- and/or protein transport-associated pro-
     teins.
    HEMBA1000012, HEMBA1000141, HEMBA1000592, HEMBA1003617,
                                                                  HEMBA1003773, HEMBA1004202,
    HEMBA1004276, HEMBA1004734, HEMBA1004847,
                                                  HEMBA1004929.
                                                                  HEMBA1004930, HEMBA1005047,
     HEMBA1005202, HEMBA1006031, HEMBA1006272.
                                                  HFMBA1006474
                                                                  HEMBA1006652, HEMBA1006914,
     HEMBA1006973, HEMBA1007224, HEMBB1000915, HEMBB1001112,
                                                                  HEMBB1001137, HEMBB1001736.
     HEMBB1001831, HEMBB1001915, MAMMA1000085, MAMMA1000734, MAMMA1001008, MAMMA1002170,
    MAMMA1002219, MAMMA1002236, MAMMA1002619, NT2RM1000661, NT2RM1000833, NT2RM2000092,
    NT2RM2000504, NT2RM2000577,
                                   NT2RM2000821, NT2RM2001201,
                                                                  NT2RM2001592, NT2RM2001613,
    NT2RM2001648, NT2RM2001730,
                                   NT2RM2001760,
                                                   NT2RM2002055.
                                                                  NT2RM4000155,
                                                                                  NT2RM4000169.
     NT2RM4000344, NT2RM4000356,
                                   NT2RM4000421, NT2RM4000712,
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                    NT2RP2001660.
                                   NT2RP2001740.
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                    NT2RP2002980.
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                                                                  NT2RP2005012,
                                                                                  NT2RP2005116,
    NT2RP2005360, NT2RP2005763, NT2RP2005784, NT2RP3000366,
     NT2RP3000759,
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                                                                                  NT2RP3002151,
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                                   NT2RP3003301,
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                                                                PLACE1000081,
                                                                               PLACE1000492,
    OVARC1001180, OVARC1001342,
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    PLACE1000863, PLACE1001092,
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    PLACE1006488, PLACE1006829.
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    [0072] The following 130 clones presumably belong to cytoskeleton-associated proteins.
    HEMBA1000156, HEMBA1000168, HEMBA1000411, HEMBA1000588, HEMBA1001043, HEMBA1001651,
                                                                HEMBA1003235, HEMBA1003581,
    HEMBA1001661, HEMBA1002102, HEMBA1002161, HEMBA1002939,
    HEMBA1004499, HEMBA1004534, HEMBA1004697, HEMBA1004929,
                                                                HEMBA1004972, HEMBA1005582,
    HEMBA1005595, HEMBA1006344, HEMBA1006737, HEMBB1001175.
                                                                HEMBB1001282, HEMBB1001562,
    HEMBB1001802, MAMMA1000824, MAMMA1001041, MAMMA1001576, MAMMA1001679, MAMMA1001735,
    MAMMA1002297, MAMMA1002351, MAMMA1002622, MAMMA1002637, MAMMA1003127, NT2RM1000850,
    NT2RM1000898, NT2RM2000030, NT2RM2000260, NT2RM2000691,
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    NT2RM4000229, NT2RM4000515, NT2RM4001217, NT2RP1000202, NT2RP1000348, NT2RP1000460,
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    NT2RP4001927, OVARC1000001, OVARC1000106.
                                                                OVARC1000520, OVARC1000679,
                                                 OVARC1000437,
    OVARC1001731, OVARC1002050, PLACE1001104, PLACE1002571,
    PLACE1002591, PLACE1002655.
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    PLACE1010960,
                                                 PLACE4000009,
                                                                THYRO1000132,
                                                                                THYRO1001405,
    PLACE2000458, PLACE3000145,
                                  PLACE3000416,
    THYRO1001458, Y79AA1000368, Y79AA1000794, Y79AA1000833, Y79AA1000962, Y79AA1002208,
    [0073] The following 54 clones presumably belong to cell division-associated and/or cell proliferation-associated
    proteins.
    HEMBA1001019, HEMBA1001595, HEMBA1002363, HEMBA1002997, HEMBA1003136, HEMBA1003369,
    HEMBA1004131, HEMBA1004354, HEMBA1005621, HEMBB1000037, HEMBB1000264, MAMMA1001768,
    MAMMA1002769, NT2RM1000354, NT2RM1000430, NT2RM1000874, NT2RM2001256, NT2RM2001743,
    NT2RM2001896, NT2RM2002145, NT2RM4000215, NT2RM4001714, NT2RP1000163, NT2RP1000333,
40
    NT2RP1000439, NT2RP2000346,
                                  NT2RP2001397.
                                                 NT2RP2002595.
                                                                 NT2RP2003177,
                                                                                NT2RP2003596.
    NT2RP2003912, NT2RP2004396.
                                  NT2RP2005037.
                                                 NT2RP2005520,
                                                                 NT2RP2005669,
                                                                                NT2RP2005835.
    NT2RP3001730, NT2RP3002081.
                                  NT2RP4000210.
                                                  NT2RP4000415.
                                                                 NT2RP4001414.
                                                                                NT2RP4001634.
                                                  PLACE1002433, PLACE1004316, PLACE1005287,
    OVARC1000013, OVARC1000937, PLACE1001383,
    PLACE1008808, PLACE1010720, PLACE1010833, Y79AA1000748, Y79AA1001236, Y79AA1001394
    [0074] The following 36 clones presumably belong to the category of embryogenesis- and/or development-associ-
    ated proteins:
    HEMBA1000518, HEMBA1001847, HEMBA1001869, HEMBA1003545, HEMBA1004973, HEMBB1002442,
    MAMMA1001837, NT2RM2001670, NT2RM4000046, NT2RM4000531, NT2RM4001140, NT2RM4001858,
    NT2RP2002078, NT2RP2004187, NT2RP2006436, NT2RP3000603,
                                                                NT2RP3000994.
                                                                                NT2RP3001580
50
    NT2RP3001708, NT2RP3003071, NT2RP3004472, NT2RP3004617,
                                                                 NT2RP4000246.
                                                                                NT2RP4001567.
    OVARC1000304, OVARC1000746, PLACE1000793,
                                                  PLACE1002532.
                                                                 PLACE1003258.
                                                                                PLACE1003625.
     PLACE1004460, PLACE1009622, PLACE4000558, THYRO1000085, Y79AA1001391, Y79AA1001692
    [0075] The following 30 clones presumably belong to cellular defense-associated proteins.
    HEMBA100005, HEMBA1000531, HEMBA1003417, HEMBA1006253, NT2RM4000354.
                                                                                NT2RM4001880,
                   NT2RP1000493,
                                   NT2RP2000006,
                                                  NT2RP2000045,
                                                                 NT2RP2000809,
                                                                                NT2RP2001536,
     NT2RP1000333,
                   NT2RP2004920.
                                   NT2RP2005037,
                                                  NT2RP3000590,
                                                                 NT2RP3001426,
                                                                                NT2RP3002062,
     NT2RP2002464.
                                                                 PLACE1006958,
     NT2RP3002785,
                   NT2RP3004262,
                                  NT2RP4001555,
                                                  NT2RP4001638,
                                                                                PLACE1008275.
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PLACE1009113, PLACE1011858, PLACE4000014, THYRO1000684, Y79AA1002139, Y79AA1002229 [0076] Although it is unclear whether or not 261 clones out of clones other than the above-mentioned clones belong to any of the above-described categories, these clones are predicted to have some functions, based on the homology search using their full-length sequences.

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HEMBA1000030, HEMBA1000307, HEMBA1000333, HEMBA1000488, HEMBA1000523, HEMBA1001197,
HEMBA1001302, HEMBA1001455, HEMBA1001675, HEMBA1001714,
                                                            HEMBA1001744, HEMBA1001967,
                             HEMBA1002458, HEMBA1002777,
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                                                                           HEMBA1003199,
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HEMBA1003615, HEMBA1003836,
                              HEMBA1004295, HEMBA1004573,
                                                            HEMBA1004604,
                                                                           HEMBA1004795,
HEMBA1005101, HEMBA1005201,
                              HEMBA1005206,
                                            HEMBA1005530,
                                                            HEMBA1005666,
                                                                           HEMBA1005990,
HEMBA1006268, HEMBA1006398,
                              HEMBA1006445,
                                             HEMBA1007174,
                                                            HEMBA1007251,
                                                                           HEMBB1000036.
                              HEMBB1001058, HEMBB1001234,
HEMBB1000144, HEMBB1000973,
                                                            HEMBB1001288,
                                                                           HEMBB1001331,
                             HEMBB1002510, HEMBB1002705,
                                                           MAMMA1000055,
                                                                           MAMMA1000625,
HEMBB1001384.
              HEMBB1002266.
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MAMMA1002972, MAMMA1003113, NT2RM1000118, NT2RM1000186,
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                              NT2RM1000883,
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                              NT2RM2001983.
                                             NT2RM2002109.
                                                            NT2RM2002142.
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                              NT2RM4000386.
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                                                            NT2RM4002075.
                                                                            NT2RM4002226.
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Y79AA1001923, Y79AA1002083, Y79AA1002307, Y79AA1002311, Y79AA1002487,
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[0077] In some cases, the predicted functions based on the partial sequences are different from those based on the full-length sequences. The reason is that a protein does not always belong solely to a single category of the above-described functional categories, and therefore, a protein may belong to two or more of the predicted functional categories. Besides, additional functions can be found for the clones classified into these functional categories by further analyses.

[0078] Since the protein encoded by clones of the invention contains full-length amino acid sequence, it is possible to analyze its biological activity, and its effect on cellular conditions such as cell proliferation and differentiation by expressing the protein as a recombinant protein using an appropriate expression system, injecting the recombinant into the cell, or raising a specific antibody against the protein.

[0079] If the protein is a secretory protein, membrane protein, or protein associated with glycoprotein, signal trans-

duction, or transcription, its biological activity can be analyzed by the methods in "Gene Transcription" (Hames B.D., and Higgins S.J. edit, (1993)), "Glycobiology" (Fukuda M., and Kobata A. edit, (1993)), "Growth Factors" (McKay I., and Leigh I. edit, (1993)), "Extracellular Matrix" (Haralson M.A., and Hassell J.R. edit, (1995), "Transcription Factors" (Latchman D.S. edit, (1993)), "Signal Transduction" (Milligans G. edit, (1992)), "Protein Phosphorylation" (Hardies G. D. edit, (1993)), and "Ion Channels" (Ashley R.H. edit, (1995) featured in "The Practical Approach Series" (IRL PRESS), or "Signal Transduction Protocols" (Kendall D.A., and Hill S.J. edit, (1995), "Glycoprotein Analysis in Biomedicine" (Hounsell E.F. edit, (1993)), featured in "Method in Molecular Biology" (Humana Press).

[0080] As to a protein associated with a disease, it is possible to perform a functional analysis as described above, but also possible to analyze correlation between the expression or the activity of the protein and a certain disease by using a specific antibody that is obtained by using expressed protein. Alternatively, it is possible to utilize the database Online Mendelian Inheritance in Man (OMIM) (http://www.ncbi.nlm.nih.gov/Omim/), which is a database of human genes and diseases, to analyze the protein.

New information is constantly being deposited in the OMIM database. Therefore, it is possible for one skilled in the art to find a new relationship between a particular disease and a gene of the present invention in the most up-to-date database.

[0081] Also, as for a secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., search of the OMIM with the following keywords resulted in the finding that the proteins are associated with many diseases (the result of the OMIM search for secrete and membrane proteins is shown below). Also, association between proteins associated to signal transduction or transcription and diseases is reported in "Transcription Factor Research-1999" (Fujii, Tamura, Kageyama, and Satake edit, (1999) Jikken-Igaku Zoukan, Vol.17, No.3), and "Gene Medicine" ((1999) Vol.3, No.2). For example, in tumors, many proteins have been shown to play a role, including secretory proteins, membrane proteins, and proteins associated with signal transduction, glycoprotein, and transcription, and also proteins associated with metabolism, cytoskeleton, and cell cycle, as described in "Tumor Biology" (Matsubara S. (1992) Syoukabou Life Science series). Thus, besides the proteins associated with diseases, many proteins described above are also potentially associated with diseases, and thus useful as a target in the medicinal industry.

[0082] The result of the OMIM search for secretory and membrane proteins is shown below, in which the keywords,

- (1) secretion protein,
- (2) membrane protein,
- (3) channel, and

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(4) extracellular matrix were used.

[0083] Shown in the search result are only the accession numbers in the OMIM. Using the number, data showing the relationship between a disease and a gene or protein can be seen. The OMIM data has been renewed everyday.

# 1) Secretion protein

268 entries found, searching for "secretion protein" 104760, 176860, 160900, 107400, 118910, 139320, 603850, 147572,176880, 600946, 603215, 157147, 600174, 151675, 170280, 179512, 179513, 138120, 179509, 246700, 179510, 600626, 179511, 600998, 109270, 601489, 154545, 179490, 185860, 603216, 122559, 601746,147290, 602672, 146770, 603062, 179508, 131230, 601591, 602434, 182590, 603166, 308230, 602534, 603489, 107470, 150390, 104610, 173120, 158106, 143890, 306900, 104610, 1046308700, 134797, 137350, 227500, 176300, 107730, 600760, 138079, 120180, 120160, 120150, 124092, 138160, 1201600, 1201600, 1201600, 1201600, 1201600, 1201600, 1201600, 12016000, 12016000, 12016000, 12016000, 120160000, 120160000, 1201600147670, 133170, 176730, 300300, 134370, 274600, 120140,162151, 158070, 152790, 120120, 106100, 300200, 192340, 190160, 138040, 147470, 147620, 173350, 147380, 152200, 152760, 157145, 153450, 264080, 113811. 600937, 600840, 188545, 202110, 600514, 186590, 603372, 136435, 137241, 252800, 214500, 207750, 138850, 136435, 1364555, 136455, 136655, 136655, 136655, 136655, 136655, 136655, 136655, 136655, 136655, 136655, 136655, 1366555, 1366555, 1366555, 1366555, 1366550, 1366550, 1366550, 1366550, 1366550, 1366550, 1366550, 1366550,139191, 142640, 138130, 189907, 603692, 600633, 603355, 107270, 600377, 147892, 232200, 600281, 232800, 602358, 137035, 601771, 601769, 253200, 601933, 118444, 600270, 120700, 600945, 603732, 147660, 600761, 172400, 600823, 600877, 130080, 171060, 107740, 307800, 602843, 130660, 152780, 124020, 601124, 601340, 601604, 601610, 171050, 312060, 232700, 300159, 142703, 600734, 125255, 168450, 123812, 188540, 147940, 188450, 600839, 182452, 188400, 182280, 176760, 263200, 600264, 188826, 252650, 601185, 162641, 137216, 162641, 188826, 1888602663, 151670, 602682, 602730, 602779, 146880, 603061, 142704, 603140, 106150, 600732, 153620, 603318,139392, 600042, 102200, 603493, 182100, 264300, 603795, 184600 2) Membrane protein

1017 entries found, searching for "membrane protein"

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                                   602014, 600442, 601109, 602481, 277900, 254210, 138247, 164920, 170280, 171050, 128100, 173910, 600884,
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                                    107269, 216550, 103320, 603489, 603551, 603767, 603799, 603842
                     [0084] There are several methods for analyzing the expression levels of genes associated with diseases. Differences
                     in gene expression levels between diseased and normal tissues are studied by the analytical methods, for example,
                     Northern hybridization and differential display. Other examples include a method with high-density cDNA filter, a method
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in gene expression levels between diseased and normal tissues are studied by the analytical methods, for example, Northern hybridization and differential display. Other examples include a method with high-density cDNA filter, a method with DNA microarray and methods with PCR amplification (Experimental Medicine, Vol.17, No. 8, 980-1056 (1999); Cell Engineering (additional volume) DNA Microarray and Advanced PCR Methods, Muramatsu & Naba (eds.), Shujunsya). The varying levels of gene expression between diseased tussues and normal tissues can be studied by any of these analytical methods. When explicit difference in the expression level is observed for a gene, it can be conclude that the gene is closely associated with a disease or disorder. Instead of diseased tissues, cultured cells can be used

for the assessment. Similarly, when gene expression is explicitly different between normal cells and cells reproducing disease-associated specific features, it can be concluded that the gene is closely associated with a disease or disorder. When the expression levels of genes are evidently varied during major cellular events (such as differentiation and apoptosis), the genes are involved in the cellular events and accordingly are candidates for disease- and/or disorder-associated genes. Further, genes exhibiting tissue-specific expression are genes playing important parts in the tissue functions and, therefore, can be candidates for genes associated with diseases and/or disorders affecting the tissues.

[0085] For example, non-enzymic protein glycation reaction is believed to be a cause for a variety of chronic diabetic complications. Accordingly, genes of which expression levels are elevated or decreased in a glycated protein-dependent manner in the endothelial cells, are associated with diabetic complications caused by glycated proteins (Diabetes 1996, 45 (Suppl. 3), S67-S72; Diabetes, 1997, 46 (Suppl. 2), S19-S25).

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The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)- $\alpha$  participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis. Many genes acting at the downstream of TNF- $\alpha$  and IL-1 $\beta$  among inflammation-associated cytokines have been previously identified. The respective stimulations are transduced through independent pathways of signaling cascade. There exists another signaling cascade for both stimulations, wherein NF- $\alpha$  B is a common transducing molecule shared by the two stimulations (J. Leukoc. Biol., 1994, 56(5): 542-547). It has also been revealed that many inflammation-associated genes, including IL-2, IL-6 and G-CSF, are varied in the expression levels thereof in response to the signal through the common pathway (Trend Genet. 1999, 15(6): 229-235). It is assumed that genes of which expression levels are varied in response to the stimulation of TNF- $\alpha$  or IL-1 $\beta$  also participate in inflammation.

[0086] Ultraviolet radiation damage has been recognized as a risk factor for skin cancers, etc. (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression tevels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with ultraviolet radiation damage of skin. In addition, genes associated with neural differentiation can be candidates for genes responsible for neurological diseases as well as candidates for genes usable for treating the diseases.

[0087] Clones exhibiting differences in the expression levels thereof can be selected by using gene expression analysis. The selection comprises, for example; analyzing cDNA clones by using high-density cDNA filter; and statistically treating the multiple signal values (signal values of radioisotope in the labeled probes or values obtained by measuring fluorescence intensities emitted from the fluorescent labels) for the respective clones by two-sample t-test, where the signal values are determined by multiple experiments of hybridization. The clones of interest are selectable based on the statistically significant differences in the signal distribution at p<0.05. However, selectable clones with significant difference in the expression levels thereof may be changed depending on the partial modification of statistical treatment. For example, the clones may be selected by conducting statistical treatment with two-sample t-test at p<0.01; or genes exhibiting more explicit differences in the expression levels thereof can be selected by performing statistical treatment with a pre-determined cut-off value for the significant signal difference. An alternative method is that the expression levels are simply compared with each other, and then, the clones of interest are selected based on the ratio of the expression levels thereof.

[0088] Clones that vary in their expression levels can also be selected by comparing the expression levels by PCR analysis, for example, by using the method of determining the band intensities representing the amounts of PCR products with ethidium bromide staining; the method of determining the values of radioisotope signals or fluorescence intensities of the PCR products when radiolabeled or fluorescent dye-labeled primers, respectively, are used in PCR amplification; or the method of determining the values of radioisotope signals or fluorescence intensities of the probes hybridized to the PCR products when radiolabeled or fluorescent dye-labeled probes, respectively, are used in the hybridization. If the expression level ratios obtained in multiple PCR experiments are constantly at least 2-fold, such a clone can be judged to vary in its expression level. When the ratios are several-fold or not less than 10-fold, the clone can be selected as a gene exhibiting the explicit difference in its expression level.

[0089] A survey of genes of which expression levels are varied specifically to the glycated protein in the endothelial cells has revealed genes with elevated expression levels, HEMBA1003958, HEMBA1004850, MAMMA1001256, MAMMA1002132, PLACE2000411 and PLACE3000119. On the other hand, a gene of which expression level is decreased specifically to the glycated protein is MAMMA1001783. These clones are genes associated with diabetes.

[0090] A survey of genes of which expression levels are varied in response to TNF-α (Tumor Necrosis Factor-alpha) in the primary cell culture of synovial tissue has revealed the following clones with elevated expression levels in the presence of TNF-α:

HEMBA1000005, HEMBA1000012, HEMBA1000020, HEMBA1000046, HEMBA1000076, HEMBA1000111, HEMBA1000168, HEMBA1000185, HEMBA1000201, HEMBA1000231, HEMBA1000243, HEMBA1000280,

HEMBA1000282, HEMBA1000304, HEMBA1000307, HEMBA1000327, HEMBA1000356, HEMBA1000376, HEMBA1000387, HEMBA1000390, HEMBA1000418, HEMBA1000460, HEMBA1000491, HEMBA1000501, HEMBA1000518, HEMBA1000519, HEMBA1000520, HEMBA1000531, HEMBA1000534, HEMBA1000542, HEMBA1000545, HEMBA1000591, HEMBA1000592, HEMBA1000594, HEMBA1000636, HEMBA1000655. HEMBA1000657, HEMBA1000673, HEMBA1000682, HEMBA1000686, HEMBA1000722, HEMBA1000726. HEMBA1000827, HEMBA1000870, HEMBA1000918, HEMBA1000971, HEMBA1000974, HEMBA1000986, HEMBA1001019, HEMBA1001043, HEMBA1001051, HEMBA1001059, HEMBA1001071, HEMBA1001060, HEMBA1001213, HEMBA1001080, HEMBA1001109, HEMBA1001140, HEMBA1001172, HEMBA1001196, HEMBA1001226, HEMBA1001281, HEMBA1001299, HEMBA1001302, HEMBA1001303, HEMBA1001323, HEMBA1001326, HEMBA1001327, HEMBA1001330, HEMBA1001351, HEMBA1001407, HEMBA1001411, 10 HEMBA1001714, HEMBA1001800, HEMBA1001446, HEMBA1001454, HEMBA1001569, HEMBA1001647, HEMBA1001921, HEMBA1001967, HEMBA1001804, HEMBA1001809, HEMBA1001888, HEMBA1001912, HEMBA1002337, HEMBA1002363, HEMBA1002084, HEMBA1002161. HEMBA1002241, HEMBA1002166, HEMBA1002542,~ HEMBA1002389, HEMBA1002458, HEMBA1002460. HEMBA1002538, HEMBA1002469, HEMBA1002547, HEMBA1002609, HEMBA1002624, 15 HEMBA1002779, HEMBA1002810, HEMBA1002816, HEMBA1002659, HEMBA1002750, HEMBA1002770, HEMBA1003021. HEMBA1003033, HEMBA1003078, HEMBA1002850, HEMBA1002863, HEMBA1002818, HEMBA1003235, HEMBA1003276, HEMBA1003229. HEMBA1003204, HEMBA1003202, HEMBA1003166, HEMBA1003403, HEMBA1003418, HEMBA1003370, HEMBA1003376. HEMBA1003296, HEMBA1003286, HEMBA1003560, HEMBA1003571, HEMBA1003591, HEMBA1003569. HEMBA1003433, HEMBA1003447, 20 HEMBA1003621. HEMBA1003662, HEMBA1003680, HEMBA1003656, HEMBA1003597, HEMBA1003598, HEMBA1003742, HEMBA1003773, HEMBA1003729, HEMBA1003733, HEMBA1003715, HEMBA1003725, HEMBA1004048, HEMBA1004074. HEMBA1004012, HEMBA1004015, HEMBA1003783, HEMBA1003950, HEMBA1004203, HEMBA1004207, HEMBA1004131, HEMBA1004202, HEMBA1004086, HEMBA1004111, HEMBA1004356, HEMBA1004366, HEMBA1004321, HEMBA1004330, HEMBA1004248, HEMBA1004274, 25 HEMBA1004507. HEMBA1004429, HEMBA1004499, HEMBA1004509, HEMBA1004405, HEMBA1004408, HEMBA1004542, HEMBA1004596, HEMBA1004604, HEMBA1004776, HEMBA1004889. HEMBA1004934, HEMBA1005274, HEMBA1004978, HEMBA1005019, HEMBA1005047, HEMBA1005206, HEMBA1005219, HEMBA1005331, HEMBA1005338, HEMBA1005394, HEMBA1005423, HEMBA1005732, HEMBA1005576, HEMBA1005746, HEMBA1006091, HEMBA1006142, HEMBA1006173, HEMBA1006198, HEMBA1006253, HEMBA1006486, HEMBA1006492, HEMBA1006268, HEMBA1006309, HEMBA1006377, HEMBA1006474, HEMBA1006659, HEMBA1006885. HEMBA1006502, HEMBA1006535, HEMBA1006579, HEMBA1006648, HEMBA1007080, HEMBA1007121, HEMBA1007194, HEMBA1006929, HEMBA1006941, HEMBA1007078, HEMBB1000044, HEMBB1000089, HEMBB1000036, HEMBA1007300, HEMBA1007301, HEMBA1007322, HEMBB1000591, HEMBB1000593. HEMBB1000215, HEMBB1000217, HEMBB1000272, HEMBB1000420, HEMBB1000975, HEMBB1000985, HEMBB1000631, HEMBB1000835, HEMBB1000887, HEMBB1000908, HEMBB1001331, HEMBB1001337, HEMBB1001014, HEMBB1001133, HEMBB1001112, HEMBB1001011, HEMBB1001394, HEMBB1001429, HEMBB1001463, HEMBB1001367, HEMBB1001384, HEMBB1001366, HEMBB1001706, HEMBB1001753, HEMBB1001797, HEMBB1001802, HEMBB1001684, HEMBB1001619, HEMBB1001910, HEMBB1001915, HEMBB1001973, HEMBB1001983, HEMBB1001812, HEMBB1001874, HEMBB1002193, HEMBB1002249, HEMBB1002329, HEMBB1002342, HEMBB1001990, HEMBB1002190, HEMBB1002442, HEMBB1002489, HEMBB1002492, HEMBB1002520, HEMBB1002371, HEMBB1002409, HEMBB1002664, HEMBB1002692, HEMBB1002697, HEMBB1002705, HEMBB1002534, HEMBB1002596, MAMMA1000092, MAMMA1000155, MAMMA1000163, MAMMA1000173, MAMMA1000175, MAMMA1000227, MAMMA1000241, MAMMA1000257, MAMMA1000264, MAMMA1000266, MAMMA1000270, MAMMA1000307, MAMMA1000410, MAMMA1000413, MAMMA1000416, MAMMA1000421, MAMMA1000472, MAMMA1000501, MAMMA1000605, MAMMA1000643, MAMMA1000670, MAMMA1000684, MAMMA1000696, MAMMA1000732, MAMMA1000752, MAMMA1000802, MAMMA1000824, MAMMA1000905, MAMMA1000921, MAMMA1000931, MAMMA1000957, MAMMA1000962, MAMMA1000998, MAMMA1001008, MAMMA1001050, MAMMA1001074, MAMMA1001078, MAMMA1001292, MAMMA1001397, MAMMA1001476, MAMMA1001743, MAMMA1001744, MAMMA1001754, MAMMA1001760, MAMMA1001785, MAMMA1001858, MAMMA1001908, MAMMA1002236, MAMMA1002267, MAMMA1002292, MAMMA1002311, MAMMA1002322, MAMMA1002359, MAMMA1002362, MAMMA1002485, MAMMA1002494, MAMMA1002597, MAMMA1002598, MAMMA1002665, MAMMA1002671, MAMMA1002684, MAMMA1002748, MAMMA1002775, MAMMA1002830, MAMMA1002858, MAMMA1002868, MAMMA1002886, MAMMA1002887, 55 MAMMA1002892, MAMMA1002909, MAMMA1002937, MAMMA1002947, MAMMA1002964, MAMMA1002970, MAMMA1003013, MAMMA1003150, NT2RM1000039, NT2RM1000062, NT2RM1000080, NT2RM1000086, NT2RM1000127, NT2RM1000132, NT2RM1000187, NT2RM1000199, NT2RM1000244, NT2RM1000256.

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                    NT2RM2000032,
                                    NT2RM2000042,
                                                    NT2RM2000101.
                                                                    NT2RM2000124,
                                                                                   NT2RM2000192,
                                    NT2RM2000363,
                                                    NT2RM2000368,
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                                                    NT2RM2002030,
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                    NT2RM4002226,
    NT2RM4002194.
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                                                                                    NT2RP2000288.
                                                    NT2RP2000097.
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                                    NT2RP1001361.
                                    NT2RP2000369,
                                                    NT2RP2000422,
                                                                    NT2RP2000842,
                                                                                    NT2RP2000965.
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                    NT2RP2000329,
                                                                                    NT2RP2001677.
     NT2RP2001245,
                    NT2RP2001440,
                                    NT2RP2001560.
                                                    NT2RP2001634,
                                                                    NT2RP2001663.
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                                    NT2RP2002312,
    NT2RP2001762,
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                                                    NT2RP2003125,
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                                                    NT2RP2003643,
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                                                                    NT2RP2005719,
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    NT2RP2005457,
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                    NT2RP3001126,
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                                                                    NT2RP3001529,
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                    NT2RP3001915,
                                    NT2RP3001969,
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                                    NT2RP3002810,
                                                     NT2RP3002876,
                                                                    NT2RP3003311,
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                                                     NT2RP4000078,
                                                                    NT2RP4000541.
     NT2RP3003672,
                    NT2RP3004209,
                                    NT2RP3004378,
                                                                                    NT2RP5003459,
                                    NT2RP4001276,
                                                     NT2RP4001507.
                                                                    NT2RP4002047,
     NT2RP4001219,
                    NT2RP4001228,
                                                    OVARC1000106.
                                                                    OVARC1000151,
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     NT2RP5003492.
                    OVARC1000085.
                                    OVARC1000087.
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                    OVARC1000440.
                                    OVARC1000564.
     OVARC1000431,
                    OVARC1000960,
                                    OVARC1000971,
                                                    OVARC1001038,
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     OVARC1000912.
    OVARC1001129,
                    OVARC1001167,
                                    OVARC1001339,
                                                    OVARC1001425.
                                                                    OVARC1001745,
                                                                                    OVARC1001762,
                                     OVARC1002044,
                                                     OVARC1002138,
                                                                    PLACE1000004,
                                                                                    PLACE1000005,
     OVARC1001766,
                     OVARC1001942,
                     PLACE1000547,
                                     PLACE1000562,
                                                     PLACE1000653,
                                                                    PLACE1001168,
                                                                                    PLACE1001311,
     PLACE1000420,
                                                                                    PLACE1002140,
                     PLACE1001920,
                                     PLACE1001983,
                                                     PLACE1002066,
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     PLACE1001377,
                                                                                    PLACE1002665,
     PLACE1002171,
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                                     PLACE1002474,
                                                     PLACE1002499,
                                                                    PLACE1002532,
                                                                                    PLACE1003783,
                                                     PLACE1003605,
                                                                    PLACE1003704,
     PLACE1003025,
                     PLACE1003145,
                                     PLACE1003361,
                                                     PLACE1004686,
                                                                    PLACE1004930,
                                                                                    PLACE1005066.
                                     PLACE1004629,
     PLACE1003885,
                     PLACE1004405,
                                                     PLACE1006143,
                                                                    PLACE1006325,
                                                                                    PLACE1006488,
                     PLACE1005630,
                                     PLACE1005876,
     PLACE1005077,
                     PLACE1006829,
                                     PLACE1007286,
                                                     PLACE1007858.
                                                                    PLACE1008201.
                                                                                    PLACE1009045,
     PLACE1006805.
                                    PLACE1010106.
                                                     PLACE1010310.
                                                                    PLACE1010622,
                                                                                    PLACE1010944,
     PLACE1009113,
                    PLACE1009621,
                     PLACE1011185.
                                     PLACE1011332,
                                                     PLACE1011635,
                                                                    PLACE1011646,
                                                                                    PLACE1011725,
     PLACE1010965,
                                                                                    PLACE3000220,
     PLACE2000014,
                     PLACE2000264,
                                     PLACE2000394,
                                                     PLACE2000419,
                                                                    PLACE3000160,
     PLACE3000254,
                     PLACE3000271,
                                     PLACE3000339,
                                                     PLACE3000341,
                                                                    PLACE3000350,
                                                                                    PLACE3000353,
                                    SKNMC1000091,
                                                     THYRO1000855,
                                                                    THYRO1001559,
                                                                                    Y79AA1000065,
     PLACE3000401,
                    PLACE4000300,
                                     Y79AA1000346,
                                                     Y79AA1000784,
                                                                     Y79AA1000833,
                                                                                    Y79AA1000968.
     Y79AA1000202.
                     Y79AA1000214.
     Y79AA1001555, Y79AA1002220
45
     [0091] On the other hand, clones with decreased expression levels in the presence of TNF\alpha are:
                                                                                    NT2RM4000290.
     HEMBA1002150,
                     HEMBB1000240,
                                     NT2RM2000469,
                                                     NT2RM2000984,
                                                                    NT2RM2001688,
                                                                                    NT2RM4001650,
                                                                    NT2RM4001611,
                                     NT2RM4001047,
                                                     NT2RM4001582.
     NT2RM4000496,
                     NT2RM4000590.
                                                     NT2RP1000243,
                                                                    NT2RP1000581,
                                                                                    NT2RP1000688,
                                     NT2RP1000174,
     NT2RM4002075,
                     NT2RM4002128.
                     NT2RP1000825,
                                     NT2RP1001185:
                                                     NT2RP1001286,
                                                                    NT2RP1001432,
                                                                                    NT2RP1001457,
     NT2RP1000767.
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                                     NT2RP2000841,
                                                     NT2RP2001813,
                                                                    NT2RP2002137,
                                                                                    NT2RP2002928,
     NT2RP2000001.
                                                                                    NT2RP2006365,
     NT2RP2003517,
                     NT2RP2003559,
                                     NT2RP2003564,
                                                     NT2RP2004933,
                                                                    NT2RP2005038,
                     NT2RP3000320,
                                     NT2RP3000484,
                                                     NT2RP3000980,
                                                                     NT2RP3001111,
                                                                                    NT2RP3001420,
     NT2RP3000072,
     NT2RP3001495.
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                                     NT2RP3002057,
                                                     NT2RP3002545,
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                                                                                    NT2RP3002799,
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                                                     NT2RP3003282,
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55
                                                     NT2RP3004262,
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                                                                                    NT2RP4000500.
     NT2RP3003385,
                     NT2RP3003870.
                                     NT2RP3004207,
                                                                    NT2RP4000989,
                                                                                    NT2RP4001442.
                     NT2RP4000787.
                                     NT2RP4000927.
                                                     NT2RP4000955.
     NT2RP4000524,
                                     NT2RP4002888,
                                                                                    PLACE1000246,
     NT2RP4001638,
                     NT2RP4001950,
                                                    NT2RP5003524,
                                                                    OVARC1001270,
     PLACE1002816.
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[0092] These are rheumatoid arthritis-associated clones. [0093] A survey of genes of which expression levels are varied in primary cultured skin fibroblast cells exposed to ultraviolet light has revealed the following clones with elevated expression levels by ultraviolet radiation: HEMBA1000542, HEMBA1001808, HEMBA1002177, HEMBA1003314, MAMMA1001874, NT2RM2001100, OVARC1001092, OVARC1001342, NT2RP2005732, NT2RP3000592, NT2RP4000657, OVARC 1000004, PLACE1002816, NT2RM4001002, NT2RM4001813, NT2RM4002266, NT2RP2001174, NT2RP2001196, PLACE1006382, THYRO1000070, NT2RP3001216, NT2RP3003464, NT2RP2005358, NT2RP3000690. THYRO1001100, Y79AA1000342 [0094] On the other hand, the expression levels of the following clones were decreased by ultraviolet radiation: HEMBA1000168, HEMBA1000231, HEMBA1000005, HEMBA1000150, HEMBA1000156, HEMBA1000158, HEMBA1000366, HEMBA1000369, HEMBA1000390, HEMBA1000333, HEMBA1000304, HEMBA1000307, HEMBA1000464, HEMBA1000469, HEMBA1000490, HEMBA1000434, HEMBA1000396, HEMBA1000418, HEMBA1000673, HEMBA1000682, HEMBA1000557. HEMBA1000657, HEMBA1000505. HEMBA1000504. HEMBA1000852, HEMBA1000870, HEMBA1000752. HEMBA1000851, HEMBA1000727, HEMBA1000686. HEMBA1001235, HEMBA1001265, HEMBA1001133, HEMBA1001121, HEMBA1001085, HEMBA1000872, HEMBA1001310, HEMBA1001323, HEMBA1001299, HEMBA1001303, HEMBA1001281, HEMBA1001289, HEMBA1001712, HEMBA1001835, HEMBA1001678, HEMBA1001640, HEMBA1001620, HEMBA1001595, HEMBA1002341, HEMBA1002419, HEMBA1002253, HEMBA1002321, HEMBA1001987, HEMBA1001950, HEMBA1002999, HEMBA1003034, HEMBA1002935, HEMBA1002728. HEMBA1002818, HEMBA1002679, HEMBA1003175, HEMBA1003202, HEMBA1003212, HEMBA1003142, 20 HEMBA1003071. HEMBA1003098. HEMBA1003447, HEMBA1003528, HEMBA1003417, HEMBA1003373, HEMBA1003276, HEMBA1003220, HEMBA1004011, HEMBA1004055, HEMBA1003885, HEMBA1003989, HEMBA1003684, HEMBA1003799, HEMBA1004272, HEMBA1004353. HEMBA1004631, HEMBA1004669, HEMBA1004133, HEMBA1004225, HEMBA1004776, HEMBA1004803, HEMBA1004816, HEMBA1004900, HEMBA1004705, HEMBA1004753, HEMBA1005101, HEMBA1005149, HEMBA1005152, HEMBA1005202, HEMBA1005047, HEMBA1005079, HEMBA1005511, HEMBA1005513, HEMBA1005518, HEMBA1005570, HEMBA1005314, HEMBA1005372, HEMBA1005853. HEMBA1005632, HEMBA1005588, HEMBA1005609, HEMBA1005577, HEMBA1005581, HEMBA1006502, HEMBA1006696. HEMBA1006031, HEMBA1006035, HEMBA1006485, HEMBA1006486, HEMBA1007301, HEMBA1007319, HEMBA1007224, HEMBA1006789, HEMBA1006796, HEMBA1007085, HEMBB1000217, HEMBB1000266, HEMBA1007341, HEMBA1007342, HEMBB1000036, HEMBB1000037, HEMBB1000399, HEMBB1000434, HEMBB1000317, HEMBB1000336, HEMBB1000369, HEMBB1000354, HEMBB1000810, HEMBB1000883, HEMBB1000673, HEMBB1000789, HEMBB1000592, HEMBB1000438, HEMBB1001424, HEMBB1001267. HEMBB1001182, HEMBB1001242, HEMBB1000887, HEMBB1001105, HEMBB1002092, HEMBB1002139, HEMBB1001531, HEMBB1001618, HEMBB1001996, HEMBB1001464, HEMBB1002550, HEMBB1002556, HEMBB1002520. HEMBB1002142, HEMBB1002190, HEMBB1002453, MAMMA1000069, MAMMA1000133, MAMMA1000055, HEMBB1002600, HEMBB1002664, MAMMA1000009, MAMMA1000585, MAMMA1000713, MAMMA1000171, MAMMA1000173, MAMMA1000287, MAMMA1000416, MAMMA1000760, MAMMA1000798, MAMMA1000831, MAMMA1000875, MAMMA1000876, MAMMA1000877, MAMMA1000906, MAMMA1000931, MAMMA1000962, MAMMA1001133, MAMMA1001139, MAMMA1001243, MAMMA1001271, MAMMA1001274, MAMMA1001298, MAMMA1001606, MAMMA1001630, MAMMA1001670, MAMMA1001743, MAMMA1001751, MAMMA1002140, MAMMA1002145, MAMMA1002158, MAMMA1002170, MAMMA1002236, MAMMA1002311, MAMMA1002498, MAMMA1002754, MAMMA1002780, MAMMA1002820, MAMMA1002843, MAMMA1002844, MAMMA1002871, MAMMA1003047, NT2RM1000037, NT2RM1000039, NT2RM1000499, NT2RM1000669, NT2RM1000746. NT2RM1000080, NT2RM1000086, NT2RM1000341, NT2RM2000260, NT2RM1000962, NT2RM2000239, NT2RM1000905, NT2RM1000885, NT2RM1000781. NT2RM2000984, NT2RM2000735, NT2RM2000821, NT2RM2000649, NT2RM2000639, NT2RM2000371, NT2RM2001194, NT2RM2001196. NT2RM2001177, NT2RM2001105, NT2RM2001065, NT2RM2001035, NT2RM2001588, NT2RM2001635, NT2RM2001648. NT2RM2001424, NT2RM2001256, NT2RM2001243, NT2RM2001730, NT2RM2001743. NT2RM2001668, NT2RM2001706. NT2RM2001727, NT2RM2001652, NT2RM2001855. NT2RM2001760, NT2RM2001771, NT2RM2001785, NT2RM2001800, 50 NT2RM2001753, NT2RM2002142. NT2RM2002049, NT2RM2002091, NT2RM2001896, NT2RM2001997. NT2RM2002030, NT2RM4000368, NT2RM4000344, NT2RM4000215, NT2RM2002145. NT2RM2002178. NT2RM2002580, NT2RM4000712, NT2RM4000515, NT2RM4000457, NT2RM4000496, NT2RM4000421. NT2RM4000425, NT2RM4000950, NT2RM4000996. NT2RM4000813, NT2RM4000820, NT2RM4000852, NT2RM4000787, NT2RM4001151, NT2RM4001187, NT2RM4001140, NT2RM4001016, NT2RM4001047, NT2RM4001054, NT2RM4001605, NT2RM4001454, NT2RM4001489, NT2RM4001258, NT2RM4001437. NT2RM4001204, NT2RM4001714, NT2RM4001715. NT2RM4001731, NT2RM4001710, NT2RM4001666, NT2RM4001611, NT2RM4001940, NT2RM4001984, NT2RM4001746, NT2RM4001856. NT2RM4001938, NT2RM4001741,

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[0095] These clones are associated with ultraviolet radiation damage.

[0096] A survey of genes of which expression levels are varied in response to the stimulation for inducing cell differentiation (stimulation using retinoic acid (RA) or using RA/inhibitor (inhibitor for cell division)) in culture cells of neural strain, NT2, revealed the following clones with elevated expression levels in the presence of RA:

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PLACE4000558, PLACE4000581, PLACE4000650, PLACE4000654.
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THYRO1000034, THYRO1000085,
                              THYRO1000092,
                                                                           THYRO1000327,
THYRO1000173, THYRO1000190,
                              THYRO1000197,
                                             THYRO1000221,
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                                                            THYRO1000585,
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                              THYRO1000394,
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THYRO1000381, THYRO1000387,
                                                            THYRO1000684,
                              THYRO1000666, THYRO1000676,
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THYRO1000637, THYRO1000658,
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                                                            THYRO1000815,
                                                                           THYRO1000865.
THYRO1000734, THYRO1000793,
                              THYRO1000796,
                                                            THYRO1001031.
                                                                           THYRO1001062,
THYRO1000916, THYRO1000934,
                              THYRO1000974,
                                             THYRO1000975.
                                             THYRO1001177,
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THYRO1001093, THYRO1001133,
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THYRO1001213, THYRO1001262,
                                             THYRO1001320,
                                                            THYRO1001322,
                              THYRO1001290,
THYRO1001406, THYRO1001426.
                              THYRO1001480,
                                             THYRO1001487,
                                                            THYRO1001537,
                                                                            THYRO1001595,
                                                            THYRO1001828,
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THYRO1001617, THYRO1001637.
                              THYRO1001706,
                                             THYRO1001772,
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Y79AA1000059,
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Y79AA1001068,
               Y79AA1001493,
Y79AA1001711,
               Y79AA1002103,
                              Y79AA1002115,
                                              Y79AA1002258,
                                                             Y79AA1002361.
                                                                            Y79AA1002407,
Y79AA1002472, Y79AA1002482
[0097] On the other hand, clones of which expression levels decreased by RA are as follows:
HEMBA1000946, HEMBA1003569, HEMBA1005570, HEMBB1000915, NT2RM1000666,
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                              NT2RM4001754, NT2RM4001905, NT2RP2001675,
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NT2RM2000594, NT2RM2001256,
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                              NT2RP3002081,
                                              NT2RP3004594,
                                                             NT2RP4001950,
               NT2RP3000980,
NT2RP2005491.
OVARC1000431, OVARC1001942, OVARC1001943, PLACE1003190, PLACE1004868,
                                                                            PLACE1005923,
PLACE1007257, PLACE1010624, Y79AA1000346
[0098] Clones of which expression levels increase by RA/inhibitor are as follows:
HEMBA1000046, HEMBA1000307, HEMBA1000434, HEMBA1000504, HEMBA1000588, HEMBA1000682,
HEMBA1000726, HEMBA1000943, HEMBA1001071, HEMBA1001094, HEMBA1001122, HEMBA1001323,
HEMBA1001361, HEMBA1001455, HEMBA1001709, HEMBA1001746, HEMBA1001869, HEMBA1002084,
HEMBA1002583, HEMBA1002628, HEMBA1002801, HEMBA1002937, HEMBA1003096, HEMBA1003142,
HEMBA1003229, HEMBA1003276, HEMBA1003309, HEMBA1003463, HEMBA1003597, HEMBA1003617,
HEMBA1003725, HEMBA1003803, HEMBA1003879, HEMBA1003989, HEMBA1004000, HEMBA1004015,
HEMBA1004024, HEMBA1004049, HEMBA1004056, HEMBA1004199, HEMBA1004248,
                                                                           HEMBA1004356.
                                             HEMBA1004770, HEMBA1004803, HEMBA1004923,
HEMBA1004554, HEMBA1004666, HEMBA1004725,
               HEMBA1004954, HEMBA1005039,
                                             HEMBA1005075, HEMBA1005113, HEMBA1005219,
HEMBA1004934,
                                             HEMBA1005367, HEMBA1005372,
               HEMBA1005251, HEMBA1005304,
                                                                            HEMBA1005403.
HEMBA1005232,
HEMBA1005410,
               HEMBA1005411, HEMBA1005548,
                                             HEMBA1005581, HEMBA1005631,
                                                                            HEMBA1005666,
                                             HEMBA1006130, HEMBA1006364,
                                                                            HEMBA1006485.
HEMBA1005755,
               HEMBA1005780, HEMBA1006067,
                                             HEMBB1000059, HEMBB1000575,
                                                                            HEMBB1000709.
HEMBA1006559,
               HEMBA1006579,
                              HEMBA1006754,
HEMBB1000822, HEMBB1000848, HEMBB1000852, HEMBB1000913, HEMBB1000985,
                                                                           HEMBB1001117.
HEMBB1001210, HEMBB1001317, HEMBB1001394, HEMBB1001443, HEMBB1001668,
                                                                            HEMBB1001695.
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HEMBB1002049, HEMBB1002254, HEMBB1002266, HEMBB1002371, HEMBB1002502, HEMBB1002614,
    HEMBB1002617, HEMBB1002692, HEMBB1002697, MAMMA1000241, MAMMA1000424, MAMMA1000616,
    MAMMA1000731, MAMMA1000824, MAMMA1000908, MAMMA1000956, MAMMA1001038, MAMMA1001091,
    MAMMA1001243, MAMMA1001815, MAMMA1001820, MAMMA1002267, MAMMA1002769, MAMMA1002871,
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                                                                                   PLACE3000353,
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                                                                   PLACE4000034,
                                                                                   PLACE4000089,
    PLACE3000362,
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                                    PLACE3000400,
    PLACE4000522, PLACE4000558,
                    THYRO1000040,
                                    THYRO1000197.
                                                   THYRO1000241.
                                                                   THYRO1000327.
                                                                                   THYRO1000394.
    SKNMC1000050.
    THYRO1000488.
                    THYRO1000501,
                                    THYRO1000585.
                                                   THYRO1000596.
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                                                                                   THYRO1000805.
                    THYRO1001133.
                                    THYRO1001134.
                                                   THYRO1001173.
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                                                                                   THYRO1001262,
    THYRO1000934.
    THYRO1001290,
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                                                                    Y79AA1000976,
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    Y79AA1001228,
                    Y79AA1001299,
                                    Y79AA1001402,
                                                    Y79AA1001585.
                                                                    Y79AA1001696.
    Y79AA1001827, Y79AA1001875, Y79AA1002027, Y79AA1002211, Y79AA1002234, Y79AA1002258
    [0099] On the other hand, clones of which expression levels decrease by RA/inhibitor are as follows:
    HEMBA1000012, HEMBA1000501, HEMBA1000946, HEMBA1003220,
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                                                                                   HEMBA1005296.
    HEMBA1003591, HEMBA1003926,
                                    HEMBA1004168,
                                                   HEMBA1004507.
    HEMBA1005528. HEMBA1005570.
                                    HEMBA1006467,
                                                   HEMBA1006486,
                                                                   HEMBA1006492.
                                                                                  HEMBA1007322.
                                                                  MAMMA1001139,
                                                                                  MAMMA1001743.
    HEMBB1000055, HEMBB1000244,
                                   HEMBB1001665,
                                                  MAMMA1000684.
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                                                   NT2RM1000666.
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    NT2RM1000257
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                                                    NT2RP2004226,
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35
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    OVARC1001942.
                    OVARC1001943.
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    PLACE1007507.
                    PLACE1008941.
                                    PLACE1010624,
    Y79AA1000346, Y79AA1001541
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[0100] These clones are also associated with neural differentiation and, therefore, are candidates for genes associated with neurological diseases.

[0101] For example, if the protein encoded by the cDNA of the present invention is a regulatory factor of cellular conditions such as growth and differentiation, it can be used for developing medicines as follows. The protein or antibody provided by the invention is injected into a certain kind of cells by microinjection. Then, using the cells, it is possible to screen low molecular weight compounds by measuring the change in the cellular conditions, or the activation or inhibition of a particular gene. The screening can be performed as follows. First, the protein is expressed and purified as recombinant. The purified protein is microinjected into cells such as various cell lines, or primary culture cells, and the cellular change such as growth and differentiation can be examined. Alternatively, the induction of genes whose expression is known to be associated with a particular change of cellular conditions may be detected by the amount of mRNA or protein. Or, the amount of intracellular molecules (low molecular weight compounds, etc.) that is changed by the function of the gene product (protein) which is known to be associated with a particular change of cellular conditions may be detected. The compounds to be screened (both low and high molecular compounds are acceptable) can be added to the culture media and assessed for their activity by measuring the change of the cellular conditions. Instead of microinjection, cell lines introduced with the gene obtained in the invention can be used for the screening. If the gene product is turn out to be associated with a particular change in the cellular conditions, the change of the product can be used as a measurement for screening. Once a compound is screened out which can activate or inhibit

the function of the protein of the invention, it can be applied for developing medicines.

[0102] If the protein encoded by the cDNA of the present invention is a secretory protein, membrane protein, or protein associated with signal transduction, glycoprotein, transcription, or diseases, it can be used in functional assays for developing medicines.

[0103] In case of a membrane protein, it is most likely to be a protein that functions as a receptor or ligand on the cell surface. Therefore, it is possible to reveal a new relationship between a ligand and receptor by screening the membrane protein of the invention based on the binding activity with the known ligand or receptor. Screening can be performed according to the known methods.

[0104] For example, a ligand against the protein of the invention can be screened in the following manner. Namely, a ligand that binds to a specific protein can be screened by a method comprising the steps of: (a) contacting a test sample with the protein of the invention or a partial peptide thereof, or cells expressing these, and (b) selecting a test sample that binds to said protein, said partial peptide, or said cells.

[0105] On the other hand, for example, screening using cells expressing the protein of the present invention that is a receptor protein can also be performed as follows. It is possible to screen receptors that is capable of binding to a specific protein by using procedures (a) attaching the sample cells to the protein of the invention or its partial peptide, and (b) selecting cells that can bind to the said protein or its partial peptide.

[0106] In a following screening as an example, first the protein of the invention is expressed, and the recombinant protein is purified. Next, the purified protein is labeled, binding assay is performed using a various cell lines or primary cultured cells, and cells that are expressing a receptor are selected (Growth and differentiation factors and their receptors, Shin-Seikagaku Jikken Kouza Vol.7 (1991) Honjyo, Arai, Taniguchi, and Muramatsu edit, p203-236, Tokyo-Kagaku-Doujin). A protein of the invention can be labeled with RI such as <sup>125</sup>I, and enzyme (alkaline phosphatase etc.). Alternatively, a protein of the invention may be used without labeling and then detected by using a labeled antibody against the protein. The cells that are selected by the above screening methods, which express a receptor of the protein of the invention, can be used for the further screening of an agonists or antagonists of the said receptor.

[0107] Once the ligand binding to the protein of the invention, the receptor of the protein of the invention or the cells expressing the receptor are obtained by screening, it is possible to screen a compound that binds to the ligand and receptor. Also it is possible to screen a compound that can inhibit both bindings (agonists or antagonists of the receptor, for example) by utilizing the binding activities.

[0108] When the protein of the invention is a receptor, the screening method comprises the steps of (a) contacting the protein of the invention or cells expressing the protein of the invention with the ligand, in the presence of a test sample, (b) detecting the binding activity between said protein or cells expressing said protein and the ligand, and (c) selecting a compound that reduces said binding activity when compared to the activity in the absence of the test sample. Furthermore, when the protein of the invention is a ligand, the screening method comprises the steps of (a) contacting the protein of the invention with its receptor or cells expressing the receptor in the presence of samples, (b) detecting the binding activity between the protein and its receptor or the cells expressing the receptor, and (c) selecting a compound that can potentially reduce the binding activity compared to the activity in the absence of the sample.

[0109] Samples to screen include cell extracts, expressed products from a gene library, synthesized low molecular compound, synthesized peptide, and natural compounds, for example, but are not construed to be listed here. A compound that is isolated by the above screening using a binding activity of the protein of the invention can also be used as a sample.

[0110] A compound isolated by the screening may be a candidate to be an agonist or an antagonist of the receptor of the protein. By utilizing an assay that monitors a change in the intracellular signaling such as phosphorylation which results from reduction of the binding between the protein and its receptor, it is possible to identify whether the obtained compound is an agonist or antagonist of the receptor. Also, the compound may be a candidate of a molecule that can inhibit the interaction between the protein and its associated proteins (including a receptor) in vivo. Such compounds can be used for developing drugs for precaution or cures of a disease with which the protein is associated.

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[0111] Secretory proteins may regulate cellular conditions such as growth and differentiation. It is possible to find out a novel factor that regulates cellular conditions by adding the secretory protein of the invention to a certain kind of cell, and performing a screening by utilizing the cellular changes in growth or differentiation, or activation of a particular gene.

[0112] The screening can be performed, for example, as follows. First, the protein of the invention is expressed and purified in a recombinant form. Then, the purified protein is added to a various kind of cell lines or primary cultured cells, and the change in the cell growth and differentiation is monitored. The induction of a particular gene that is known to be involved in a certain cellular change is detected by the amounts of mRNA and protein. Alternatively, the amount of an intracellular molecule (low-molecular-weight compounds, etc.) that is changed by the function of a gene product (protein) that is known to function in a certain cellular change is used for the detection.

[0113] Once the screening reveals that the protein of the invention can regulate cellular conditions or the functions, it is possible to apply the protein as a pharmaceutical and diagnostic medicine for associated diseases by itself or by

altering a part of it into an appropriate composition.

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[0114] As is above described for membrane proteins, the secretory protein provided by the invention may be used to explore a novel ligand-receptor interaction using a screening based on the binding activity to a known ligand or receptor. A similar method can be used to identify an agonist or antagonist. The resulting compounds obtained by the methods can be a candidate of a compound that can inhibit the interaction between the protein of the invention and an interacting molecule (including a receptor). The compounds may be able to use as a preventive, therapeutic, and diagnostic medicine for the diseases, in which the protein may play a certain role.

[0115] Proteins associated with signal transduction or transcription may be a factor that affects a certain protein or gene in response to intracellular/extracellular stimuli. It is possible to find out a novel factor that can affect a protein or gene by expressing the protein provided by the invention in a certain types of cells, and performing a screening utilizing the activation of a certain intracellular protein or gene.

[0116] The screening may be performed as follows. First, a transformed cell line expressing the protein is obtained. Then, the transformed cell line and the untransformed original cell line are compared for the changes in the expression of a certain gene by detecting the amount of its mRNA or protein. Alternatively, the amount of an intracellular molecule (low molecular weight compounds) that is changed by the function of a certain gene product (protein) may be used for the detection. Furthermore, the change of the expression of a certain gene can be detected by introducing a fusion gene that comprises a regulatory region of the gene and a marker gene (luciferase, beta-galactosidase, etc.) into a cell, expressing the protein provided by the invention into the cell, and estimating the activity of a marker gene product (protein).

[0117] If the protein or gene of the invention is associated with diseases, it is possible to screen a gene or compound that can regulate its expression and/or activity either directly or indirectly by utilizing the protein of the present invention.

[0118] For example, the protein of the invention is expressed and purified as a recombinant protein. Then, the protein or gene that interacts with the protein of the invention is purified, and screened based on the binding. Alternatively, the screening can be performed by adding with a compound of a candidate of the inhibitor added in advance and monitoring the change of binding activity. In another method, a transcription regulatory region locating in the 5'-upstream of the gene encoding the protein of the invention that is capable of regulating the expression of other genes is obtained, and fused with a marker gene. The fusion is introduced into a cell, and the cell is added with compounds to explore a regulatory factor of the expression of the said gene.

[0119] The compound obtained by the screening can be used for developing pharmaceutical and diagnostic medicines for the diseases with which the protein of the present invention is associated. Similarly, if the regulatory factor obtained in the screening is turn out to be a protein, compounds that can newly affect the expression or activity of the protein may be used as a medicine for the diseases with which the protein of the invention is associated.

[0120] If the protein of the invention has an enzymatic activity, regardless as to whether it is a secretory protein, membrane protein, or proteins associated with signal transduction, glycoprotein, transcription, or diseases, a screening may be performed by adding a compound to the protein of the invention and monitoring the change of the compound. The enzymatic activity may also be utilized to screen a compound that can inhibit the activity of the protein.

[0121] In a screening given as an example, the protein of the invention is expressed and the recombinant protein is purified. Then, compounds are contacted with the purified protein, and the amount of the compound and the reaction products is examined. Alternatively, compounds that are candidates of an inhibitor are pretreated, then a compound (substrate) that can react with the purified protein is added, and the amount of the substrate and the reaction products is examined.

[0122] The compounds obtained in the screening may be used as a medicine for diseases with which the protein of the invention is associated. Also they can be applied for tests that examine whether the protein of the invention functions normally *in vivo*.

[0123] Whether the secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein of the present invention is a novel protein associated with diseases or not is determined in another method than described above, by obtaining a specific antibody against the protein of the invention, and examining the relationship between the expression or activity of the protein and a certain disease. In an alternative way, it may be analyzed referred to the methods in "Molecular Diagnosis of Genetic Diseases" (Elles R. edit, (1996) in the series of "Method in Molecular Biology" (Humana Press).

[0124] Proteins associated with diseases are targets of screening as mentioned, and thus are very useful in developing drugs which regulate their expression and activity. Also, the proteins are useful in the medicinal industry as a diagnostic marker of the associated disease or a target of gene therapy.

[0125] Compounds isolated as mentioned above can be administered patients as it is, or after formulated into a pharmaceutical composition according to the known methods. For example, a pharmaceutically acceptable carrier or vehicle, specifically sterilized water, saline, plant oil, emulsifier, or suspending agent can be mixed with the compounds appropriately. The pharmaceutical compositions can be administered to patients by a method known to those skilled in the art, such as intraarterial, intravenous, or subcutaneous injections. The dosage may vary depending on the weight

or age of a patient, or the method of administration, but those skilled in the art can choose an appropriate dosage properly. If the compound is encoded by DNA, the DNA can be cloned into a vector for gene therapy, and used for gene therapy. The dosage of the DNA and the method of its administration may vary depending on the weight or age of a patient, or the symptoms, but those skilled in the art can choose properly.

[0126] The present invention further relates to databases comprising at least a sequence of polynucleotide and/or protein, or a medium recorded in such databases, selected from the sequence data of the nucleotide and/or the amino acids indicated in Table 350 and Table 351.

The term "database" means a set of accumulated information as machine-searchable and readable information of nucleotide sequence. The databases of the present invention comprise at least one of the novel nucleotide sequences of polynucleotides provided by the present invention. The databases of the present invention can consist of only the sequence data of the novel polynucleotides provided by the present invention or can comprise other information on nucleotide sequences of known full-length cDNAs or ESTs. The databases of the present invention can be comprised of not only the information on the nucleotide sequences but also the information on the gene functions revealed by the present invention. Additional information such as names of DNA clones carrying the full-length cDNAs can be recorded or linked together with the sequence data in the databases.

[0127] The database of the present invention is useful for gaining complete gene sequence information from partial sequence information of a gene of interest. The database of the present invention comprises nucleotide sequence information of full-length cDNAs. Consequently, by comparing the information in this database with the nucleotide sequence of a partial gene fragment yielded by differential display method or subtraction method, the information on the full-length nucleotide sequence of interest can be gained from the sequence of the partial fragment as a starting clue. [0128] The sequence information of the full-length cDNAs constituting the database of the present invention contains not only the information on the complete sequences but also extra information on expression frequency of the genes as well as homology of the genes to known genes and known proteins. Thus the extra information facilitates rapid functional analyses of partial gene fragments. Further, the information on human genes is accumulated in the database of the present invention, and therefore, the database is useful for isolating a human homologue of a gene originating from other species. The human homologue can be isolated based on the nucleotide sequence of the gene from the original species.

[0129] At present, information on a wide variety of gene fragments can be obtained by differential display method and subtraction method. In general, these gene fragments are utilized as tools for isolating the full-length sequences thereof. When the gene fragment corresponds to an already-known gene, the full-length sequence is easily obtained by comparing the partial sequence with the information in known databases. However, when there exists no information corresponding to the partial sequence of interest in the known databases, cDNA cloning should be carried out for the full-length cDNA. It is often difficult to obtain the full-length nucleotide sequence using the partial sequence information as an initial clue. If the full-length of the gene is not available, the amino acid sequence of the protein encoded by the gene remains unidentified. Thus the database of the present invention can contribute to the identification of full-length cDNAs corresponding to gene fragments, which cannot be revealed by using databases of known genes.

[0130] The present invention has provided 5602 novel full-length cDNA clones, and primers for synthesizing the cDNA. As has not yet proceeded the isolation of full-length cDNA within the human, the invention has great significance. The full-length cDNA clones contain the translation initiation site, and thus provide a useful information for analysis of protein functions.

[0131] The cDNA clones are assumed to encode proteins such as secretory proteins, membrane proteins, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., which have important functions in vivo, and also predicted to be associated with many diseases. The genes and proteins associated with diseases are useful for developing a diagnostic marker or medicines for regulation of their expression and activity, or as a target of gene therapy.

[0132] The invention is illustrated more specifically with reference to the following examples, but is not to be construed as being limited thereto.

# **EXAMPLE 1**

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Construction of a cDNA library by the oligo-capping method.

[0133] The NT-2 neuron progenitor cells (Stratagene), a teratocarcinoma cell line from human embryo testis, which can differentiate into neurons by the treatment with retinoic acid were used.

The NT-2 cells were cultured according to the manufacturer's instructions as follows.

- (1) NT-2 cells were cultured without induction by retinoic acid treatment (NT2RM1, NT2RM2, NT2RM4).
- (2) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 48 hours (NT2RP1).

(3) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 2 weeks (NT2RP2, NT2RP3, NT2RP4, NT2RP5).

[0134] Also, the human neuroblastoma cell line SK-N-MC (ATCC HTB-10) (SKNMC1), and human retinoblastoma cell line Y79 (ATCC HTB-18) (Y79AA1) were cultured according to the culture conditions described in the ATCC catalogue (http://www.atcc.org/). The cells were harvested separately, and mRNA was extracted from each cell by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)+RNA was purified from the mRNA using oligo-dT cellulose. [0135] Similarly, human placenta (PLACE1, PLACE2, PLACE3, PLACE4), human ovary cancer tissue (OVARC1), tissues from human embryo at 10 weeks, which is enriched with head (HEMBA1), or body (HEMBB1), human mammary gland (MAMMA1), human thyroid gland (THYRO1), and primary cultured cells of human blood vessel endothelium (VESEN1) were used to extract mRNA by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)+RNA was purified from the mRNA using oligo-dT cellulose.

[0136] Each poly(A)+RNA was used to construct a cDNA library by the oligo-capping method (Maruyama M. and Sugano S. (1994) Gene, 138: 171-174). Using the Oligo-cap linker (SEQ ID NO: 10464) and the Oligo-dT primer (SEQ ID NO: 10465), bacterial alkaline phosphatase (BAP) treatment, tobacco acid phosphatase (TAP) treatment, RNA ligation, the first strand cDNA synthesis, and removal of RNA were performed as described in the reference (Suzuki and Kanno (1996) Protein Nucleic acid and Enzyme, 41: 197-201; Suzuki Y. et al. (1997) Gene, 200: 149-156). Next, 5'- and 3'-PCR primers (SEQ ID NO: 10466, and 10467, respectively) were used for performing PCR to convert the cDNA into double stranded cDNA, which was then digested with Sfil. Then, the DrallI-cleaved pUC19FL3 vector (Figure 1; for NT2RM1, and NT2RP1), or the DrallI-cleaved pME18SFL3 (Figure 1) (GenBank AB009864, expression vector; for NT2RM2, NT2RM4, NT2RP2, NT2RP3, NT2RP4, NT2RP5, SKNMC1, Y79AA1, PLACE1, PLACE2, PLACE3, PLACE4, OVARC1, HEMBA1, HEMBB1, MAMMA1, THYRO1, and VESEN1) was used for cloning the cDNA clones was analyzed with a DNA sequencer (ABI PRISM 377, PE Biosystems) after sequencing reactions were performed with the DNA sequencing reagents (Dye Terminator Cycle Sequencing FS Ready Reaction Kit, dRhodamine Terminator Cycle Sequencing FS Ready Reaction Kit, or BigDye Terminator Cycle Sequencing FS Ready Reaction Kit, PE Biosystems), according to the instructions. The data were compiled into a database.

[0137] The full-length-enriched cDNA libraries except those for NT2RM1 and NT2RP1 were constructed using eukaryotic expression vector pME18SFL3. The vector contains SRα promoter and SV40 small t intron in the upstream of the cloning site, and SV40 polyA added signal sequence site in the downstream. As the cloning site of pME18SFL3 has asymmetrical DrallI sites, and the ends of cDNA fragments contain Sfil sites complementary to the DrallI sites, the cloned cDNA fragments can be inserted into the downstream of the SRα promoter unidirectionally. Therefore, clones containing full-length cDNA can be expressed transiently by introducing the obtained plasmid directly into COS cells. Thus, the clones can be analyzed very easily in terms of the proteins that are the gene products of the clones, or in terms of the biological activities of the proteins.

[0138] Herein, the cDNA libraries and the name of each clone are related as shown in Table 3. Therein, "xxxxxx" represents the clone number of six digits. Thus, the sequences are named by the library name, the clone number plus F- for the 5'-end, or R- for the 3'-end.

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Table 3

5	library: clone	5'-end sequence	3'-end sequence
10	NT2RM1: NT2RM1xxxxxx NT2RP1:	F-NT2RM1xxxxxx	
	NT2RP1xxxxxx NT2RM2:		
15	NT2RM2xxxxxx NT2RM4:		
20	NIZKM4XXXXX	F-NT2RM4xxxxxx	N-NI ZNM4XXXXX
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	NT2RP2:		
	NT2RP2xxxxxx	F-NT2RP2xxxxxx	R-NT2RP2xxxxxx
5	NT2RP3:		
v	NT2RP3xxxxxx	F-NT2RP3xxxxxx	R-NT2RP3xxxxxx
	NT2RP4:		
	NT2RP4xxxxxx	F-NT2RP4xxxxxx	R-NT2RP4xxxxxx
10	NT2RP5:		
	NT2RP5xxxxxx	F-NT2RP5xxxxxx	R-NT2RP5xxxxxx
	SKNMC1:		
	SKNMC1xxxxxx	F-SKNMC1xxxxxx	R-SKNMC1xxxxxx
15	Y79AA1:		
	Y79AA1xxxxxx	F-Y79AA1xxxxxx	R-Y79AA1xxxxxx
	PLACE1:		
20	PLACE1xxxxxx	F-PLACE1xxxxxx	R-PLACE1xxxxxx
20	PLACE2:		
	PLACE2xxxxxx	F-PLACE2xxxxxx	R-PLACE2xxxxxx
	PLACE3:		
25	PLACE3xxxxxx	F-PLACE3xxxxxx	R-PLACE3xxxxxx
	PLACE4:		•
	PLACE4xxxxxx	F-PLACE4xxxxxx	R-PLACE4xxxxxx
	OVARC1:		·
30	OVARC1xxxxxx	F-0VARC1xxxxxx	R-0VARC1xxxxxx
	HEMBA1:		
	HEMBA1xxxxxx	F-HEMBA1xxxxxx	R-HEMBA1xxxxxx
25	HEMBB1:		
35	HEMBB1xxxxxx	F-HEMBB1xxxxxx	R-HEMBB1xxxxxx
	MAMMA1:		
	MAMMA1xxxxxx	F-MAMMA1xxxxxx	R-MAMMA1xxxxxx
40	THYRO1:		
	THYR01xxxxxx	F-THYR01xxxxxx	R-THYR01xxxxxx
	VESEN1:	e vearus	D VEOCHI
	VESEN1xxxxxx	F-VESEN1xxxxxx	R-VESEN1xxxxxx
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# **EXAMPLE 2**

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Estimation of the fullness ratio at the 5'-ends of the clones contained in the cDNA libraries constructed by the oligocapping method.

[0139] The fullness ratio at the 5'-end sequence of the 59,823 clones in the human cDNA libraries constructed by the oligo-capping method was determined as follows. Of all the clones whose 5'-end sequences were found in those of known human mRNA in the public database, a clone was judged to be "full-length", if it had a longer 5'-end sequence than that of the known human mRNA, or, even though the 5'-end sequence was shorter, if it contained the translation initiation codon. A clone which did not contain the translation initiation codon was judged to be "not-full-length". The fullness ratio ((the number of full-length clones)/(the number of full-length and not-full-length clones)) at the 5'-end of the cDNA clones from each library was determined by comparing with the known human mRNA. As a result, the fullness

ratio of the 5'-ends was 63.5%. The result indicates that the fullness ratio at the 5'-end sequence was extremely high.

#### **EXAMPLE 3**

5 Assessment of the fullness ratio of the 5'-end of the cDNA by the ATGpr and the ESTiMateFL.

[0140] The ATGpr, developed by Salamov A.A., Nishikawa T., and Swindells M.B. in the Helix Research Institute, is a program for prediction of the translation initiation codon based on the characteristics of the sequences in the vicinity of the ATG codon. The results are shown with expectations (also described as ATGpr1 below) that an ATG is a true initiation codon (0.05-0.94) (can be described as ATGpr1). When the program was applied to the 5'-end sequences of the clones from the cDNA library that was obtained by the oligo-capping method and that had 65% fullness ratio, the sensitivity and specificity of estimation of a full-length clone (clone containing the N-terminal end of ORF) were improved to 82-83% by selecting only clones having the ATGpr1 score 0.6 or higher.

[0141] Furthermore, the program was used to assess the fullness of 18,959 clones in the human fibraries obtained here, which have 5'-ends matched to a known human mRNA.

[0142] Briefly, the maximal ATGpr1 score of the clones was determined, and then their 5'-end sequence was compared with the known human mRNA to estimate whether the clone is full-length or not. The result is shown in Table 4. [0143] Based on the knowledge that known mRNAs, in general, are highly expressed in the cell, similar estimation was performed with genes having a low number in the EST hit, which represent relatively low abundant mRNAs, and the result is shown in Table 5.

[0144] In the table, the number of full-length clones indicate that of clones containing the N-terminal end of ORF, and so does the number of not-full-length clones that of clones without the N-terminal end of ORF. The fullness ratio represents (the number of full-length clones)/(the number of full-length clones plus the number of not-full-length clones).

Table 4

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having a matched 5'-end with that of a known mRNA.

number of full-length fullness ratio maximal ATGpr1 score number of (full-length clones plus not-full-length clones clones) 9,346 83.5% >=0.70 11,193 78.9% 10,549 >=0.50 13.369 73.2% 11,340 >=0.30 15,489 67.9% 17,394 11,811 >=0.15 63.5% 12,046 >=0.00 18,959

Table 5

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of the clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having 5 EST hits or less among the clones having a matched 5'-end with that of a known mRNA.

fullness ratio number of (full-length number of full-length maximal ATGpr1 score clones plus not-full-length clones clones) 69.0% 2,801 1,934 >=0.70 65.0% 3,683 2,393 >=0.50 >=0.30 4.683 2.707 57.8% 52.0% >=0.15 5.559 2.890 49.8% >=0.00 6,113 3.013

[0145] Next, the ESTiMateFL was used for the estimation. The ESTiMateFL, developed by Nishikawa and Ota in the Helix Research Institute, is a method for the selection of a clone with high fullness ratio by comparing with the 5'-end or 3'-end sequences of ESTs in the public database.

[0146] By the method, a cDNA clone is judged to be most likely not to be full-length if there exist any ESTs which

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have longer 5'-end or 3'-end sequences than the clone. The method is systematized for high throughput analysis. A clone is judged to be full-length if the clone has a longer 5'-end sequence than ESTs in the public database. Even if a clone has a shorter 5'-end, the clone is judged to be full-length if the difference in length is within 50 bases, and otherwise judged not to be full-length, for convenience.

[0147] In case of the clones whose 5'-end sequence is matching with the known mRNA, 80% of the clones judged to be full-length by comparing with ESTs was also judged to be full-length by comparing with the known mRNA. Also, 80% of the clones judged to be not full-length by comparing with ESTs was also judged to be not full-length by comparing with the known mRNA.

[0148] The precision of the estimation by comparing with ESTs is improved with increasing number of ESTs to be compared. However, in case that a limited number of ESTs are available, the reliability becomes low. Thus, the method is effective in excluding clones with high probability of being not-full-length, from the cDNA clones that is synthesized by the oligo-capping method and that have the 5'-end sequences with about 60 % fullness ratio. In particular, the ESTiMateFL is efficiently used to estimate the fullness ratio at the 3'-end sequence of cDNA of a human unknown mRNA which has a significant number of EST deposits in the public database.

[0149] The 18,959 clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, were estimated by using the ATGpr and ESTiMateFL. Briefly, the 5'-end sequence of the respective clone was analyzed to obtain the maximal ATGpr1 score, and compared with the ORF of the known human mRNA that matches with it to determine whether the clone is full-length or not. Then, the 5'-end sequence of the respective clone was analyzed by the ESTiMateFL to judge whether the clone is full-length or not. Specifically, the 5'-end sequences of the 18,959 clones were compared with those of ESTs by the ESTiMateFL and the clones other than those that are not full-length were selected. Then, the selected clones were used to analyze the relationship between the ATGpr and the fullness ratio. The result was summarized in Table 6. Also, among the selected, the clones in which the number of the EST hit is not more than 5 were selected and analyzed. The result was summarized in Table 7, which represents the result of the analysis of mRNA with relatively low abundance.

[0150] Therein, the number of being full-length, the number of being not full-length, and the fullness ratio indicate the number of the clones that contain the N-terminus of the ORF, the number of the clones that do not contain the N-terminus of the ORF, and (the number of being full-length)/(the number of being full-length) plus (the number of being not full-length), respectively.

Table 6

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequence in the clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, and also other than those being not full-length according to the comparison with ESTs.

maximal ATGpr1 score number of (full-length number of full-length fullness ratio clones plus not-full-length clones clones) >=0.70 9,068 92.1% 8,349 >=0.50 10.345 90.1% 9,318 >=0.3011,425 9,964 87.2% >=0.15 12,254 10,335 84.3% >=0.00 12,785 10,484 82.0%

Table 7

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio	
>=0.70	1,959	1,510	77.1%	٦
>=0.50	2,469	1,821	73.8%	
>=0.30	2,975	2,046	68.8%	
>=0.15	3,368	2,164	64.3%	

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#### Table 7 (continued)

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.00	3,661	2,226	60.8%

[0151] The 19,226 clones, isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to that of a known human mRNA were estimated by the ATGpr2, and the correlation between the score and the fullness ratio was estimated. Specifically, the maximal ATGpr2 score of the clones identical to a known human mRNA was determined, and then their fullness ratio was estimated by comparing the 5'-ends with ORF of known human mRNA. The result was shown in Table 8.

#### Table 8

Maximal ATGpr2 score and fullness ratio of the 5'-end sequence of the clones, which are isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA.

maximal ATGpr2 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.30	10,748	8,031	74.7%
>=0.15	16,383	11,226	68.5%
>=0.00	19,226	12,285	63.9%

[0152] According to the above results, it was found that, in case of using clones isolated from human cDNA libraries constructed by the oligo-capping method, the fullness ratio of the clones that have low score in the ATGpr can be improved by estimating their 5'-end sequence using the combination of the ATGpr and the ESTiMateFL. Therefore, the method was applied to select a cDNA clone with high fullness ratio.

## 35 EXAMPLE 4

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Clustering of the 5'-end and 3'-end sequences of cDNA clones.

[0153] The 5'-end and 3'-end sequences of cDNA clones were obtained, and clustered separately. The single pass data of the nucleotide sequence of the 5'-end and 3'-end was subject to the BLAST search between the sequence data of all the clones synthesized in example 1, and the clones considered to be originating from the same gene were put together into a group. If the 5'-end of a clone contains the consensus sequence of 300 bases or more with identity 95% or more, or the 3'-end contains the consensus sequence of 200 bases or more and having identity 90% or more, the clones were put in the same group.

[0154] The groups of the 5'-end sequence and the 3'-end sequence were further clustered so as that the groups from the same clone can be in the same group (cluster).

## **EXAMPLE 5**

- 50 Characterization of the cloned sequence.
  - [0155] The data of the 5'-end sequence of the cloned sequence was characterized by the following way:
- (1) examining whether it is identical to those of mRNA from human and other species (including authorized sequences) and human EST by the BLAST homology search of the GenBank,
  - (2) examining whether it has longer 5'-end than those of human mRNA and human EST,
  - (3) determining the scores in the ATGpr1 and ATGpr2 programs of all the initiation codons in the 5'-end sequence, and

- (4) determining the number of the human EST clone(s) that is judged to be identical by the BLAST homology search of the GenBank.
- [0156] The data of the 3'-end sequence of the cloned sequence was characterized by the above (1) and (4).
- [0157] These characterized data were used for the final selection of the clones.

#### **EXAMPLE 6**

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Identity to the human mRNA and human EST, and comparison of the length of the 5'-end.

[0158] The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of mRNA from human or other species when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 94% or more. The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of human EST when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 90% or more.

[0159] The sequence of the clone was judged to be full-length in comparison with human mRNA when the sequence has longer 5'-end, or it contains the translation initiation site. The sequence of the clone was judged to be full-length in comparison with human EST in the database when the sequence has longer 5'-end, or while it has shorter end, the difference in length between the two sequences is 50 bases or less. The other clones were judged to be not full-length.

**EXAMPLE 7** 

Prediction of the fullness ratio by the ATGpr.

25 [0160] The score in the ATGpr1 is the expectation to be full-length based on calculations, and the higher score reflects the higher probability to be full-length as shown in Example 3. The maximal ATGpr1 score and the maximal ATGpr2 score represent the score obtained with all the initiation codons contained in the 5'-end sequence of the cloned sequence, and were used for the characterization.

#### 30 EXAMPLE 8

Prediction of the novelty using the number of the identical ESTs by the homology search.

[0161] For both the 5'-end and 3'-end sequences of the clones, the number of the identical ESTs was determined by the homology search on the GenBank. Human ESTs were judged to be identical when the EST has a sequence of 200 nucleotides or more with 90% or more matching with the 5'-end sequence. The number of the identical ESTs were used for characterization and as an index of novelty. The clone having not identical sequence at the 5'-end and 3'-end sequences to those of mRNA as well as those of ESTs is a gene encoding a novel protein. Similarly, a clone having either the 5'-end or the 3'-end sequences, which has low number of the identical ESTs, is judged to be a gene encoding a novel protein.

## **EXAMPLE 9**

Characterization of clusters.

[0162] The clusters of the groups of the 5'-end and 3'-end sequences were characterized according to the following criteria.

(1) Whether it is identical to the mRNA sequences from human or other species (including authorized sequences), or human ESTs by the BLAST search of the GenBank.

A cluster containing at least one sequence of all the 5'-end and 3'-end sequences, which is identical to one of the mRNA sequences, was regarded to be the same cluster of the mRNA sequence.

(2) Whether it has longer 5'-end than human mRNA sequence and human ESTs.

When all the 5'-end sequences contained in a cluster are judged to be not full-length compared with the mRNA sequences and human ESTs, the cluster was regarded as being not full-length.

(3) The scores in the ATGpr1 and ATGpr2 using all the initiation codons contained in the 5'-end sequences. The maximal ATGpr1 score among those of all the 5'-end sequences in a cluster was determined as the ATGpr1 score of the cluster. The ATGpr2 score of the cluster was also determined in the same way.

(4) The number of the identical human ESTs determined by the BLAST search of the GenBank.

[0163] The maximum number was determined in the numbers of ESTs identical to each of 5'-end sequences contained in a cluster. The number of the ESTs identical to the 5'-end sequences in the cluster was defined as the maximum number. The number of the ESTs identical to the 3'-end sequences in a cluster was determined in the same way.

## **EXAMPLE 10**

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Methods for selection of the clusters by the characteristics.

[0164] Data obtained by the characterization described above was used to discard the clusters that are identical to any mRNA sequence from human and other species (including authorized sequences), or those clusters that are not full-length. From the rest of the clusters, the clusters that fulfill any of the following conditions were selected.

- (a) A cluster in which the number of the identical ESTs for the 5'-end sequence is 20 or less, and the ATGpr1 score exceeds 0.3.
- (b) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for both the 5'-end sequence and the 3'-end sequence is 5 or less, and multiple clones are contained.
- (c) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is 0, and the number of the identical ESTs for the 3'-end sequence is not less than 1.
- (d) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is not less than 1 and not more than 5, and the number of the identical ESTs for the 3'-end sequence is 0.

[0165] The clusters selected by (a) contain at least one clone that is novel and having high fullness ratio. The clusters selected by (b), (c), and (d) contain at least one clone that is novel and having low fullness ratio, but is still full-length.

#### **EXAMPLE 11**

Methods for selection of clones from clusters.

[0166] In the clusters comprising a single clone, the clone was selected.

[0167] In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score higher than 0.3, a clone with the highest score was selected.

[0168] In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score not more than 0.3, a clone with the highest ATGpr2 score was selected, if the score was higher than 0.3.

[0169] In the clusters comprising multiple clones, in which the clones have the scores not more than 0.3 in both the ATGpr1 and the ATGpr2, a clone with the highest scores in both the ATGpr1 and ATGpr2 was selected.

[0170] In the clusters comprising multiple clones, in which the above selection by the ATGpr score was not applicable, selected was a clone having longer 5'-end by assembling the 5'-end sequence, 3'-end sequence, and human ESTs. For assembling, the Sequencher<sup>TM</sup> (Hitachi Soft Engineering) was used. When even the selection by assembling failed, all the clones were judged to be full-length.

[0171] As a result, 3690 clones were the clones that have the maximal ATGpr1 score higher than 0.3. On the other hand, 477 clones were the clones that have the maximal ATGpr1 score not more than 0.3, and the maximal ATGpr2 score higher than 0.3. The number of the clones having the highest scores in both the ATGpr1 and ATGpr2, while the scores were not more than 0.3, were 97. The number of the clones which were not selected by the ATGpr scores, but were selected by assembling the 5'-end sequence, 3'-end sequence, and human ESTs, were 117. The clones that have the score in both the ATGpr1 and ATGpr2 not more than 0.3, but were selected because the cluster comprises a single clone, were 1166. In the clones, at least either of the 5'-end or 3'-end sequence was not identical to any of human ESTs. Some clones were selected because the cluster comprises a single clone, or by assembling, in which there is no ATG codon (9 clones: HEMBA1001960, HEMBA106569, HEMBB1001454, NT2PR2002839, NT2RP2005325, NT2RP2006323, PLACE1004506, PLACE1005526, and THYRO1001177). The sequences that do not contain the ATG codon were considered to be corresponding to the 5'-UTR. Although the clones do not have the scores in the ATGpr1 and ATGpr2, the clones were yet judged to be full-length according to the fullness ratio, as shown in Table 4, 5, 6, 7, and 8. The above clones that were finally judged to be full-length were classified into 11 groups according to the following criteria.

Group (1): 1516 clones

Among the 3690 clones having the maximal ATGpr1 score higher than 0.3, the following 1516 clones were having

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high fullness ratio and a novel clone, in which at least either of the 5'-end or 3'-end sequence, or both of them were
    not identical to any of human ESTs.
    HEMBA1000046, HEMBA1000050, HEMBA1000129, HEMBA1000150, HEMBA1000158, HEMBA1000193 .
    HEMBA1000201, HEMBA1000216, HEMBA1000227, HEMBA1000288, HEMBA1000290, HEMBA100030 3,
    HEMBA1000304, HEMBA1000369, HEMBA1000392, HEMBA1000396, HEMBA1000488, HEMBA10005 05,
    HEMBA1000508, HEMBA1000534, HEMBA1000542, HEMBA1000594, HEMBA1000637, HEMBA1000 657.
    HEMBA1000752, HEMBA1000867, HEMBA1000869, HEMBA1000872, HEMBA1000910, HEMBA100 0918,
    HEMBA1000919, HEMBA1000942, HEMBA1000968, HEMBA1000975, HEMBA1000986, HEMBA10 01022,
    HEMBA1001043, HEMBA1001052, HEMBA1001080, HEMBA1001085, HEMBA1001109, HEMBA1 001140,
    HEMBA1001174, HEMBA1001235, HEMBA1001286, HEMBA1001302, HEMBA1001398, HEMBA 1001407,
    HEMBA1001415, HEMBA1001446, HEMBA1001476, HEMBA1001497, HEMBA1001510, HEMB A1001533,
    HEMBA1001570, HEMBA1001581, HEMBA1001635, HEMBA1001640, HEMBA1001647, HEM BA1001661,
    HEMBA1001731, HEMBA1001744, HEMBA1001746, HEMBA1001800, HEMBA1001815, HE MBA1001822,
    HEMBA1001866, HEMBA1001896, HEMBA1001910, HEMBA1001987,
    HEMBA1002018, HEMBA1002035, HEMBA1002049, HEMBA1002092, HEMBA1002119, HEMBA1002125.
    HEMBA1002161, HEMBA1002177, HEMBA1002189, HEMBA1002191, HEMBA1002199, HEMBA100222 9,
    HEMBA1002237, HEMBA1002265, HEMBA1002363, HEMBA1002417, HEMBA1002419, HEMBA10024 30,
    HEMBA1002439, HEMBA1002477, HEMBA1002503, HEMBA1002508, HEMBA1002515, HEMBA1002 547,
    HEMBA1002688, HEMBA1002703, HEMBA1002746, HEMBA1002750, HEMBA1002850, HEMBA100 2973,
20
    HEMBA1003021, HEMBA1003067, HEMBA1003077, HEMBA1003078, HEMBA1003079, HEMBA10 03117,
    HEMBA1003129, HEMBA1003175, HEMBA1003199, HEMBA1003235, HEMBA1003250, HEMBA1 003257,
    HEMBA1003291, HEMBA1003322, HEMBA1003327, HEMBA1003370, HEMBA1003380, HEMBA 1003395,
    HEMBA1003402, HEMBA1003461, HEMBA1003480, HEMBA1003538, HEMBA1003545, HEMB A1003556,
    HEMBA1003581, HEMBA1003621, HEMBA1003645, HEMBA1003667, HEMBA1003720, HEM BA1003760.
    HEMBA1003799, HEMBA1003807, HEMBA1003827, HEMBA1003836, HEMBA1003866, HE MBA1003879.
    HEMBA1003880, HEMBA1003985, HEMBA1003989,
    HEMBA1004011, HEMBA1004048, HEMBA1004056, HEMBA1004074, HEMBA1004133, HEMBA1004146,
    HEMBA1004150, HEMBA1004199, HEMBA1004200, HEMBA1004238, HEMBA1004246, HEMBA100427 5,
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    HEMBA1004372, HEMBA1004389, HEMBA1004479, HEMBA1004499, HEMBA1004507, HEMBA1004 542,
    HEMBA1004554, HEMBA1004573, HEMBA1004596, HEMBA1004632, HEMBA1004697, HEMBA100 4705,
    HEMBA1004709, HEMBA1004711, HEMBA1004736, HEMBA1004751, HEMBA1004752, HEMBA10 04753,
    HEMBA1004756, HEMBA1004758, HEMBA1004768, HEMBA1004771, HEMBA1004795, HEMBA1 004806,
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### Group (2): 377 clones

. 35

Among the 3690 clones, the following 377 clones were full-length, and a novel clone, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

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## 50 Group (3): 1797 clones

Among the 3690 clones, the following 1797 clones were full-length, and novel clones, in which the number of the identical human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).

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	Y79AA1002487,					

#### Group (4): 453 clones

20

Among the 1857 clones having the maximal ATGpr1 score not more than 0.3 (including the 9 clones whose 5'-end sequence does not contain the ATG codon), the following 453 clones were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.

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## Group (5): 24 clones

The following 24 clones having low score in the ATGpr1 were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA1000622, HEMBA1000749, HEMBA1000876, HEMBA1001226, HEMBA1001391, HEMBA1002742, HEMBA1003908, HEMBA1004000, HEMBB1000336, MAMMA1000356, MAMMA1000883, MAMMA100159 0, NT2RP2000289, NT2RP2001467, PLACE1002853, PLACE1003420, PLACE1004836, PLACE10049 13, PLACE1006795, PLACE1008181, PLACE1008715, PLACE4000344, THYRO1000129, THYRO1001 321,

### Group (6): 65 clones

The following 65 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.

HEMBA1000604, HEMBA1000673, HEMBA1001024, HEMBA1001026, HEMBA1001734, HEMBA1001784,

HEMBA1001808, HEMBA1003902, HEMBA1004164, HEMBA1004909, HEMBA1005232, HEMBA100557 7, HEMBA10066461, HEMBA1006695, HEMBB10011119, HEMBB1001337, HEMBB1001536, HEMBB10018 68, HEMBB1002045, HEMBB1002579, MAMMA1000444, MAMMA1000761, MAMMA1000943, MAMMA1001 820, MAMMA1002360, NT2RM4000366, NT2RM4001856, NT2RM4002390, NT2RP2000108, NT2RP2000257, NT2RP2001506, NT2RP2002047, NT2RP2002066, NT2RP2002475, NT2RP2004400, NT2RP2004587, NT2RP200528 9, NT2RP2005694, NT2RP3001898, NT2RP3003264, NT2RP3003433, NT2RP3003842, OVARC10012 40, PLACE1001323, PLACE1002227, PLACE1002500, PLACE1002604, PLACE1002772, PLACE1003 478, PLACE1004681, PLACE1005108, PLACE1005932, PLACE1006318, PLACE1006368, PLACE100 6506, PLACE1006904, PLACE1007557, PLACE1007877, PLACE1009048, PLACE1011109, PLACE10 11643, PLACE4000548, THYRO1000279, Y79AA1000410, Y79AA1002103,

#### Group (7): 32 clones

The following 32 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA1000251, HEMBA1001803, HEMBA1001918, HEMBA1002257, HEMBA1003064, HEMBA1003714, HEMBA1004405, HEMBA1005508, HEMBB1000054, HEMBB1001142, MAMMA1000175, MAMMA100116 2, MAMMA1002972, NT2RM4000425, NT2RP2004512, NT2RP2005531, NT2RP2005942, NT2RP20065 54, NT2RP3001007, NT2RP3001318, OVARC1000017, OVARC1000068, OVARC1000486, PLACE1001 705, PLACE1002319, PLACE1007743, PLACE1007829, PLACE1008630, PLACE1009925, PLACE101 1492, PLACE1011749, THYPO1000793

PLACE1011749, THYRO1000793, The following 117 clones, selected by assembling the sequence of the other clones in the same cluster and human ESTs, have high fullness ratio. The clones were classified into the following group (8) and (9). HEMBA1001323, HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1002538, HEMBA1003309, HEMBA1003939, HEMBA1004015, HEMBA1004295, HEMBA100467 2. HEMBA1004865, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1006779, HEMBA10072 88, HEMBB1000218, HEMBB1000272, HEMBB1000399, HEMBB1000491, HEMBB1000996, HEMBB1001 114, HEMBB1001850, HEMBB1002015, MAMMA1000287, MAMMA1001683, MAMMA1001686, MAMMA100 2612. NT2RM2000609, NT2RM4002438, NT2RM4002567, NT2RP2000270, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002736, NT2RP2002753, NT2RP2003456, NT2RP200372 7. NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2004580, NT2RP20052 93, NT2RP2005476, NT2RP2005753, NT2RP2005815, NT2RP2005841, NT2RP2005857, NT2RP2006 393, NT2RP2006467, NT2RP3000109, NT2RP3000449, NT2RP3001245, NT2RP3001634, NT2RP300 2056, NT2RP3002810, NT2RP3002955, NT2RP3003032, NT2RP3003138, NT2RP3003500, NT2RP30 03819, NT2RP4000078, NT2RP4000515, NT2RP4000517, NT2RP4001407, NT2RP4001889, NT2RP4 002905, OVARC1000071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691, PLACE1001746, PLACE1001748, PLACE1001845, PLACE1002066, PLACE1003373, PLACE100390 0, PLACE1004118, PLACE1004256, PLACE1004284, PLACE1004336, PLACE1004506, PLACE10049 34. PLACE1005077, PLACE1005111, PLACE1005409, PLACE1005730, PLACE1006076, PLACE1006 360, PLACE1006470, PLACE1006760, PLACE1006867, PLACE1007045, PLACE1007111, PLACE100 7807, PLACE1008080, PLACE1008244, PLACE1008369, PLACE1008405, PLACE1008426, PLACE10 08621. PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, PLACE3000276, THYRO1 000805,

## Group (8): 36 clones

THYRO1001365, THYRO1001673, Y79AA1001848,

Among the 117 clones described above, the following 36 clones were novel clones, in which at least either of the 5'-end or 3'-end sequence, or both of them were not identical to any of human ESTs.

HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1003939, HEMBA1004295, HEMBA1004865, HEMBA1006779, HEMBB1000218, HEMBB1000491, HEMBB100099 6. HEMBB1001114, MAMMA1000287, MAMMA1001686, MAMMA1002612, NT2RP2002736, NT2RP20037 27, NT2RP2004580, NT2RP2005841, NT2RP2006393, PLACE1002066, PLACE1003373, PLACE1003 900, PLACE1004118, PLACE1004336, PLACE1004506, PLACE1004934, PLACE1005409, PLACE100 5730, PLACE1006470, PLACE1008080, PLACE3000276, THYRO1000805, THYRO1001365, THYRO10 01673.

## Group (9): 81 clones

Among the 117 clones described above, the following 81 clones were the clones in which the number of the identical

human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).

HEMBA1001323, HEMBA1002538, HEMBA1003309, HEMBA1004015, HEMBA1004672, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1007288, HEMBB1000272, HEMBB1000399, HEMBB100185 0, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP200124, NT2RP2002753, NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2005753, NT2RP2005753, NT2RP2005815, NT2RP2005857, NT2RP2006467, NT2RP200505753, NT2RP3001634, NT2RP3002056, NT2RP3002810, NT2RP3002955, NT2RP3003138, NT2RP3003500, NT2RP3003819, NT2RP400078, NT2RP4000515, NT2RP4000517, NT2RP4001407, NT2RP4001889, NT2RP4002905,

OVARC100071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691, PLACE1001746, PLACE1001748, PLACE1001845, PLACE1004256, PLACE1004284, PLACE100507 7, PLACE1005111, PLACE1006076, PLACE1006360, PLACE1006760, PLACE1006867, PLACE10070 45, PLACE1007111, PLACE1007807, PLACE1008244, PLACE1008369, PLACE1008405, PLACE1008 426, PLACE1008621, PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, Y79AA100 1848,

### Group (10): 938 clones

20

The following 938 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 4. The clones were novel clones, in which at least the 5'-end sequence was not identical to any of human ESTs.

HEMBA100042, HEMBA1000213, HEMBA1000264, HEMBA1000355, HEMBA1000366, HEMBA1000411, HEMBA1000422, HEMBA1000428, HEMBA1000434, HEMBA1000442, HEMBA1000464, HEMBA100054 0, HEMBA1000722, HEMBA1000747, HEMBA1000769, HEMBA1000773, HEMBA1000774, HEMBA1000852, HEMBA1000852, HEMBA1000870, HEMBA1000908, HEMBA1000908, HEMBA1000908, HEMBA1000908, HEMBA1001327, HEMBA1001375, HEMBA1001094, HEMBA1001094, HEMBA1001121, HEMBA1001172, HEMBA1001515, HEMBA1001557, HEMBA1001566, HEMBA1001585, HEMBA1001589, HEMBA1001589, HEMBA1001960, HEMBA1001964, HEMBA1001991, HEMBA1002003, HEMBA1002008, HEMBA1002003, HEMBA1002008, HEMBA1002039, HEMBA1002100, HEMBA1002113, HEMBA1002139,

HEMBA1002144, HEMBA1002153, HEMBA1002226, HEMBA1002253, HEMBA1002270, HEMBA100232 1, HEMBA1002328, HEMBA1002337, HEMBA1002381, HEMBA1002486, HEMBA1002552, HEMBA10025 55, HEMBA1002558, HEMBA1002561, HEMBA1002590, HEMBA1002628, HEMBA1002629, HEMBA1002651, HEMBA1002666, HEMBA1002678, HEMBA1002712, HEMBA1002716, HEMBA1002730, HEMBA1002748, HEMBA1002780, HEMBA1002801, HEMBA1002826, HEMBA1002833, HEMBA10 02886, HEMBA1003083, HEMBA1003086, HEMBA1003098, HEMBA1003133, HEMBA1 003197, HEMBA1003314, HEMBA1003376, HEMBA1003330, HEMBA1003348, HEMBA1003376, HEMBA1003384, HEMBA1003595, HEMBA1003595, HEMBA1003758, HEMBA1003758, HEMBA1003773, HEMBA1003784, HEMBA1003784, HEMBA1003976, HEMBA1003976,

HEMBA1004012, HEMBA1004045, HEMBA1004138, HEMBA1004241, HEMBA1004272, HEMBA1004306, HEMBA1004312, HEMBA1004334, HEMBA1004354, HEMBA1004356, HEMBA1004366, HEMBA100439 4, HEMBA1004396, HEMBA1004429, HEMBA1004460, HEMBA1004461, HEMBA1004482, HEMBA10045 06, HEMBA1004538, HEMBA1004586, HEMBA1004617, HEMBA1004629, HEMBA1004631, HEMBA1004 666, HEMBA1004670, HEMBA1004733, HEMBA1004816, HEMBA1004820, HEMBA1004918, HEMBA100 4956, HEMBA1005275, HEMBA1005293, HEMBA1005304, HEMBA1005315, HEMBA1005372, HEMBA1 005389, HEMBA1005570, HEMBA1005500, HEMBA1005506, HEMBA1005511, HEMBA1005520, HEMBA 1005568, HEMBA1005570, HEMBA1005705, HEMBA1005627, HEMBA1005631, HEMBA1005632, HEMBA1005670, HEMBA1005509, HEMBA1005705, HEMBA1005765, HEMBA1005780, HEMBA1005853, HEMBA1005909, HEMBA1005934, HEMBA1005962, HEMBA1005999,

HEMBA1006002, HEMBA1006005, HEMBA1006042, HEMBA1006067, HEMBA1006090, HEMBA1006142, HEMBA1006268, HEMBA1006334, HEMBA1006359, HEMBA1006416, HEMBA1006419, HEMBA100642 1, HEMBA1006446, HEMBA1006471, HEMBA1006486, HEMBA1006489, HEMBA1006546, HEMBA10065 69,

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HEMBA1006595, HEMBA1006617, HEMBA1006635, HEMBA1006648, HEMBA1006665, HEMBA1006 821.
HEMBA1006849, HEMBA1006865, HEMBA1006929, HEMBA1007045, HEMBA1007051, HEMBA100 7073,
HEMBA1007113, HEMBA1007256, HEMBA1007273, HEMBA1007322, HEMBA1007347, HEMBB1000039,
HEMBB1000044, HEMBB1000059, HEMBB1000089, HEMBB1000099, HEMBB1000113, HEMBB1000141,
HEMBB1000250, HEMBB1000264, HEMBB1000274, HEMBB1000284, HEMBB100030 7, HEMBB1000312,
HEMBB1000317, HEMBB1000337, HEMBB1000341, HEMBB1000343, HEMBB10003 69, HEMBB1000376,
HEMBB1000434, HEMBB1000441, HEMBB1000455, HEMBB1000472, HEMBB1000 518, HEMBB1000530,
HEMBB1000564, HEMBB1000575, HEMBB1000586, HEMBB1000598, HEMBB100 0637, HEMBB1000638,
HEMBB1000643, HEMBB1000665, HEMBB1000726, HEMBB1000738, HEMBB10 00770, HEMBB1000794,
HEMBB1000821, HEMBB1000822, HEMBB1000883, HEMBB1000888, HEMBB1 000890, HEMBB1000910,
HEMBB1000959, HEMBB1000981, HEMBB1001037, HEMBB1001051, HEMBB 1001060, HEMBB1001133,
HEMBB1001208, HEMBB1001209, HEMBB1001218, HEMBB1001221, HEMB B1001249, HEMBB1001254,
HEMBB1001267, HEMBB1001282, HEMBB1001302, HEMBB1001304, HEM BB1001335, HEMBB1001348,
HEMBB1001356, HEMBB1001366, HEMBB1001380, HEMBB1001443, HE MBB1001463, HEMBB1001464,
HEMBB1001521, HEMBB1001527, HEMBB1001537, HEMBB1001555, HEMBB1001586, HEMBB1001619,
HEMBB1001630, HEMBB1001637, HEMBB1001641, HEMBB1001706, HEMBB1001735, HEMBB100175 3,
HEMBB1001762, HEMBB1001802, HEMBB1001880, HEMBB1001899, HEMBB1001921, HEMBB10019 30,
HEMBB1001952, HEMBB1001953, HEMBB1001973, HEMBB1001988, HEMBB1002002, HEMBB1002 005,
HEMBB1002009, HEMBB1002042, HEMBB1002049, HEMBB1002069, HEMBB1002115, HEMBB100 2139.
HEMBB1002189, HEMBB1002232, HEMBB1002254, HEMBB1002255, HEMBB1002280, HEMBB10 02306,
HEMBB1002364, HEMBB1002371, HEMBB1002381, HEMBB1002453, HEMBB1002477, HEMBB1 002509,
HEMBB1002531, HEMBB1002601, HEMBB1002603, HEMBB1002610, HEMBB1002613, HEMBB 1002623,
HEMBB1002635, HEMBB1002664, HEMBB1002677, HEMBB1002683, HEMBB1002686, HEMB B1002699,
HEMBB1002712.
MAMMA100009, MAMMA1000045, MAMMA1000103, MAMMA1000133, MAMMA1000134, MAMMA1000155,
MAMMA1000241, MAMMA1000254, MAMMA1000264, MAMMA1000266, MAMMA1000340, MAMMA100038 5,
MAMMA1000423, MAMMA1000424, MAMMA1000446, MAMMA1000478, MAMMA1000501, MAMMA10005 16.
MAMMA1000565, MAMMA1000585, MAMMA1000605, MAMMA1000616, MAMMA1000621, MAMMA1000 643,
MAMMA1000669, MAMMA1000696, MAMMA1000707, MAMMA1000714, MAMMA1000718, MAMMA100 0732.
MAMMA1000733, MAMMA1000839, MAMMA1000845, MAMMA1000855, MAMMA1000863, MAMMA10 00867.
MAMMA1000880, MAMMA1000905, MAMMA1000942, MAMMA1000957, MAMMA1000987, MAMMA1 000998,
MAMMA1001003, MAMMA1001024, MAMMA1001030, MAMMA1001035, MAMMA1001050, MAMMA 1001067,
MAMMA1001082, MAMMA1001092, MAMMA1001145, MAMMA1001161, MAMMA1001203, MAMM A1001243,
MAMMA1001268, MAMMA1001280, MAMMA1001298, MAMMA1001322, MAMMA1001324, MAM MA1001383,
MAMMA1001408, MAMMA1001435, MAMMA1001442, MAMMA1001446,
MAMMA1001501, MAMMA1001575, MAMMA1001606, MAMMA1001671, MAMMA1001715, MAMMA1001740,
MAMMA1001788, MAMMA1001818, MAMMA1001854, MAMMA1001878, MAMMA1001931, MAMMA100199 2,
MAMMA1002032, MAMMA1002082, MAMMA1002084, MAMMA1002108, MAMMA1002118, MAMMA10021 40,
MAMMA1002215, MAMMA1002267, MAMMA1002298, MAMMA1002299, MAMMA1002310, MAMMA1002 322.
MAMMA1002339, MAMMA1002347, MAMMA1002352, MAMMA1002361, MAMMA1002392, MAMMA100 2417,
MAMMA1002428, MAMMA1002446, MAMMA1002475, MAMMA1002480, MAMMA1002545, MAMMA10 02556,
MAMMA1002566, MAMMA1002597, MAMMA1002603, MAMMA1002623, MAMMA1002625, MAMMA1 002629,
MAMMA1002636, MAMMA1002662, MAMMA1002698, MAMMA1002708, MAMMA1002721, MAMMA 1002744,
MAMMA1002754, MAMMA1002758, MAMMA1002780, MAMMA1002833, MAMMA1002835, MAMM A1002838.
MAMMA1002844, MAMMA1002858, MAMMA1002880, MAMMA1002887, MAMMA1002908, MAM MA1002930.
MAMMA1002970, MAMMA1002982, MAMMA1003003, MAMMA1003019, MAMMA1003026, MA MMA1003031.
MAMMA1003040, MAMMA1003055,
NT2RM4000046, \quad NT2RM4000085, \quad NT2RM4000200, \quad NT2RM4000244, \quad NT2RM4000368, \quad NT2RM4000979 \ , \quad NT2RM4000085, \quad NT2RM4000979 \ , \quad NT2RM40000979 \ , \quad NT2RM4000979 \ , \quad NT
NT2RM4001016, NT2RM4001519, NT2RM4001557, NT2RM4001605, NT2RM4001810, NT2RM400185 8,
NT2RM4001930, NT2RM4002067, NT2RM4002278, NT2RM4002281, NT2RM4002287, NT2RM40023 83,
NT2RM4002479, NT2RM4002504, NT2RM4002532, NT2RP2000027, NT2RP2000098, NT2RP2000 258,
NT2RP2000327, NT2RP2000337, NT2RP2000459, NT2RP2000498, NT2RP2000987, NT2RP200 1277,
 NT2RP2001312, NT2RP2001423, NT2RP2001445, NT2RP2001449, NT2RP2001677, NT2RP20 01699,
NT2RP2001720, NT2RP2001740, NT2RP2001762, NT2RP2001813, NT2RP2001869, NT2RP2 001943,
NT2RP2002032, NT2RP2002070, NT2RP2002105, NT2RP2002137, NT2RP2002192, NT2RP 2002256,
NT2RP2002259, NT2RP2002394, NT2RP2002439, NT2RP2002457, NT2RP2002504, NT2R P2002606,
 NT2RP2002727, NT2RP2002740, NT2RP2002752, NT2RP2002839, NT2RP2002987, NT2RP2003129,
 NT2RP2003161, NT2RP2003164, NT2RP2003230, NT2RP2003339, NT2RP2003499, NT2RP2003522,
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    NT2RP2004339, NT2RP2004347, NT2RP2004425, NT2RP2004490, NT2RP2004 675, NT2RP2004709,
    NT2RP2004967, NT2RP2005020, NT2RP2005031, NT2RP2005108, NT2RP200 5254, NT2RP2005325,
    NT2RP2005336, NT2RP2005344, NT2RP2005354, NT2RP2005360, NT2RP20 05491, NT2RP2005501,
    NT2RP2005581, NT2RP2005645, NT2RP2005651, NT2RP2005701, NT2RP2 005719, NT2RP2005741,
    NT2RP2005908, NT2RP2005980, NT2RP2006098, NT2RP2006103, NT2RP 2006166, NT2RP2006184,
    NT2RP2006320, NT2RP2006321, NT2RP2006323, NT2RP2006534, NT2R P2006598,
    NT2RP3000186, NT2RP3000235, NT2RP3000247, NT2RP3000433, NT2RP3000441, NT2RP3000578.
    NT2RP3000584, NT2RP3000622, NT2RP3000628, NT2RP3000685, NT2RP3000742, NT2RP300081 5,
10
    NT2RP3000875, NT2RP3000904, NT2RP3001115, NT2RP3001232, NT2RP3001374, NT2RP30014 59,
    NT2RP3001527, NT2RP3001607, NT2RP3001752, NT2RP3001782, NT2RP3001926, NT2RP3001 989,
    NT2RP3002002, NT2RP3002166, NT2RP3002352, NT2RP3002687, NT2RP3002713, NT2RP300 2770,
    NT2RP3002799, NT2RP3002978, NT2RP3003059, NT2RP3003121, NT2RP3003464, NT2RP30 03572.
    NT2RP3003746, NT2RP3003799, NT2RP3003989, NT2RP3004016, NT2RP3004095, NT2RP3 004110,
    NT2RP3004145, NT2RP3004148, NT2RP3004215, NT2RP3004399, NT2RP3004475, NT2RP 3004503,
    NT2RP4000321, NT2RP4000519, NT2RP4000996,
    OVARC1000133, OVARC1000198, OVARC1000240, OVARC1000427, OVARC1000431, OVARC1000533 ,
    OVARC1000543, OVARC1000573, OVARC1000578, OVARC1000678, OVARC1000679, OVARC100070 0.
    OVARC1000769, OVARC1000802, OVARC1000891, OVARC1000897, OVARC1000936, OVARC10009 37,
    OVARC1000960, OVARC1000971, OVARC1001000, OVARC1001062, OVARC1001072, OVARC1001 085,
    OVARC1001118, OVARC1001129, OVARC1001169, OVARC1001261, OVARC1001268, OVARC100 1282,
    OVARC1001330, OVARC1001339, OVARC1001342, OVARC1001357, OVARC1001442, OVARC10 01480,
    OVARC1001542, OVARC1001547, OVARC1001615, OVARC1001726, OVARC1001795, OVARC1 001900,
    OVARC1001901, OVARC1001911, OVARC1001989, OVARC1002044, OVARC1002066, OVARC 1002143,
    PLACE1000031, PLACE1000040, PLACE1000078, PLACE1000094, PLACE1000214, PLACE1000332,
    PLACE1000421, PLACE1000424, PLACE1000481, PLACE1000749, PLACE1000841, PLACE100093 1,
    PLACE1000979, PLACE1001015, PLACE1001076, PLACE1001118, PLACE1001136, PLACE10012 79,
    PLACE1001384, PLACE1001468, PLACE1001502, PLACE1001534, PLACE1001603, PLACE1001 611,
    PLACE1001640, PLACE1001716, PLACE1001720, PLACE1001745, PLACE1001799, PLACE100 1821,
    PLACE1001897, PLACE1002052, PLACE1002150, PLACE1002256, PLACE1002493, PLACE10 02514,
    PLACE1002578, PLACE1002591, PLACE1002598, PLACE1002768, PLACE1002782, PLACE1 002881,
    PLACE1002962, PLACE1003153, PLACE1003258, PLACE1003343, PLACE1003375, PLACE 1003401,
    PLACE1003519, PLACE1003521, PLACE1003528, PLACE1003575, PLACE1003583, PLAC E1003584,
    PLACE1003638, PLACE1003760, PLACE1003768, PLACE1003833, PLACE1003850, PLA CE1003858,
    PLACE1003932,
    PLACE1004242, PLACE1004257, PLACE1004425, PLACE1004467, PLACE1004471, PLACE1004518,
    PLACE1004658, PLACE1004693, PLACE1004813, PLACE1004827, PLACE1004838, PLACE100484 0,
    PLACE1004900, PLACE1004972, PLACE1005146, PLACE1005162, PLACE1005453, PLACE10054 71,
    PLACE1005477, PLACE1005502, PLACE1005526, PLACE1005528, PLACE1005574, PLACE1005 584.
    PLACE1005611, PLACE1005802, PLACE1005884, PLACE1005898, PLACE1005968, PLACE100 6002,
    PLACE1006129, PLACE1006143, PLACE1006187, PLACE1006205, PLACE1006223, PLACE10 06382,
    PLACE1006521, PLACE1006534, PLACE1006540, PLACE1006617, PLACE1006640, PLACE1 006779,
    PLACE1006792, PLACE1006800, PLACE1006805, PLACE1006815, PLACE1006860, PLACE 1006878,
    PLACE1007097, PLACE1007132, PLACE1007140, PLACE1007276, PLACE1007286, PLAC E1007525,
    PLACE1007677, PLACE1007737, PLACE1007866,
    PLACE1008045, PLACE1008111, PLACE1008201, PLACE1008231, PLACE1008392, PLACE1008424,
    PLACE1008437, PLACE1008455, PLACE1008625, PLACE1008748, PLACE1008757, PLACE100879 8,
    PLACE1008807, PLACE1008920, PLACE1008947, PLACE1009150, PLACE1009246, PLACE10093 08,
    PLACE1009410, PLACE1009477, PLACE1009613, PLACE1009639, PLACE1009879, PLACE1009 888,
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    PLACE1011503, PLACE1011641, PLACE1011649, PLACE1011650, PLACE1011729, PLACE 1011778,
    PLACE1011874, PLACE1011962, PLACE1011995,
    PLACE2000103, PLACE2000113, PLACE2000115, PLACE2000132, PLACE2000140, PLACE2000176,
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### Group (11): 228 clones

The following 228 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 7. The clones were a novel clone, in which at least the 3'-end sequence was not identical to any of human ESTs.

HEMBA1000111, HEMBA1000243, HEMBA1000338, HEMBA1000390, HEMBA1000418, HEMBA1000459, HEMBA1000518, HEMBA1000791, HEMBA1000791, HEMBA1001077, HEMBA1001077, HEMBA10010979, HEMBA1001123, HEMBA1001294, HEMBA1001299, HEMBA1001442, HEMBA1001861, HEMBA1001979, HEMBA1002592, HEMBA1002679, HEMBA1002944, HEMBA1002046, HEMBA1003034, HEMBA1003034, HEMBA1003571, HEMBA1003579, HEMBA1003598, HEMBA1003630, HEMBA1003034, HEMBA1003038, HEMBA1003039, HEMBA1003039, HEMBA1004049, HEMBA1004132, HEMBA1003039, HEMBA1005039, HEMBA1005152, HEMBA1005159, HEMBA1005359, HEMBA1005443, HEMBA1005497, HEMBA1005552, HEMBA1005359, HEMBA1005408, HEMBA1005329, HEMBA1005329, HEMBA1006328, HEMBA1006328, HEMBA1006338, HEMBA1006338, HEMBA1006338, HEMBA1006338, HEMBA1006338, HEMBA1006338, HEMBA1006338, HEMBA1006334, HEMBA1006338, HEMBA1006338, HEMBA1006334, HEMBA1006780, HEMBA1006921, HEMBA1006938, HEMBA1007017, HEMBA1007341,

HEMBB100050, HEMBB1000420, HEMBB1000487, HEMBB1000490, HEMBB1000523, HEMBB1000684, HEMBB1000840, HEMBB1000876, HEMBB1000913, HEMBB1000915, HEMBB1000917, HEMBB100106 3, HEMBB1001102, HEMBB1001177, HEMBB1001253, HEMBB1001271, HEMBB1001454, HEMBB10015 00, HEMBB1001588, HEMBB1001618, HEMBB1001704, HEMBB1001717, HEMBB1001756, HEMBB1001 867, HEMBB1001967, HEMBB1002043, HEMBB1002094, HEMBB1002190, HEMBB1002520, HEMBB100 2556, HEMBB1002590.

MAMMA1000431, MAMMA1000483, MAMMA1000559, MAMMA1000723, MAMMA1000744, MAMMA100075 2, MAMMA1000782, MAMMA1000802, MAMMA1000851, MAMMA1000877, MAMMA1000962, MAMMA10010 74, MAMMA1001133, MAMMA1001191, MAMMA1001206, MAMMA1001249, MAMMA1001330, MAMMA1001 452, MAMMA1001502, MAMMA1001663, MAMMA1001806, MAMMA1001836, MAMMA1001880, MAMMA1001908, MAMMA1001963, MAMMA1002058, MAMMA1002155, MAMMA1002158, MAMMA1002230, MAMMA1002282, MAMMA1002293, MAMMA1002332, MAMMA1002359, MAMMA1002494, MAMMA1 002573,

MAMMA1000117, MAMMA1000227, MAMMA1000270, MAMMA1000302, MAMMA1000348, MAMMA1000413,

MAMMA1002618, MAMMA1002622, MAMMA1002646, MAMMA1002748, MAMMA1002871, MAMMA 1002892, MAMMA1003140, NT2RM4000327, NT2RM4000532, NT2RM4001414, NT2RM4001776, NT2RM4002499, NT2RP2000076, NT2RP2000644, NT2RP2000731, NT2RP2001106, NT2RP200167, 8

NT2RP2000523, NT2RP2000603, NT2RP2000644, NT2RP2000731, NT2RP2001196, NT2RP200167 8, NT2RP2001926, NT2RP2003206, NT2RP2003704, NT2RP2003912, NT2RP2004396, NT2RP20046 81, NT2RP3000512, NT2RP3000542, NT2RP3000736, NT2RP3000865, NT2RP3002057, NT2RP3003 384, NT2RP3004334, NT2RP3004349, NT2RP3004527, NT2RP4000049, OVARC1000058, OVARC100 0191, OVARC1000302, OVARC1001051, OVARC1001117, OVARC1001668, OVARC1001745, OVARC10 01812, OVARC1001928,

PLACE100583, PLACE1001272, PLACE1001414, PLACE1002537, PLACE1003205, PLACE1003361, PLACE1003516, PLACE1003864, PLACE1004384, PLACE1004491, PLACE1004793, PLACE100498 5, PLACE1005085, PLACE1005666, PLACE1005845, PLACE1006164, PLACE1006262, PLACE10067 54, PLACE1007852, PLACE1008331, PLACE1008643, PLACE1009050, PLACE1009155, PLACE1009 172, PLACE1009183, PLACE1009200, PLACE1009398, PLACE1009595, PLACE1010702, PLACE101 0891, PLACE1011165, PLACE1011203, PLACE1011375, PLACE1011576, PLACE1011762, PLACE10 11964, PLACE2000136, PLACE2000264, PLACE2000305, PLACE2000379, PLACE3000220, PLACE4 000250,

THYRO1000092,THYRO1000471, THYRO1000852, THYRO1001534, THYRO1001595, THYRO 1001745.

# EXAMPLE 12

Homology search using the 5'-end and 3'-end sequences of the selected clones.

[0172] The 5' -end sequences of the selected 5547 clones were used for the homology search of the SwissProt, and both the 5' -end 3' -end sequences were used for the search of the GenBank and UniGene (ref. the result of the search of the SwissProt, GenBank (except ESTs and STSs), and UniGene (Human) was attached).

[0173] Each search result is shown in the last part of this SPECIFICATION by arranging each item in the following

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	5'-end sequence	3'-end sequence
Swiss-Prot	Homology search result 1	
GenBank	Homology search result 2	Homology search result 3
UniGene	Homology search result 4	Homology search result 5

[0174] According to the top hit data, at least 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth. Among the clones predicted belonging to any of the categories, 1001 clones were estimated to have a relatively high homology with the known proteins or genes in the same category. In addition, 429 clones were estimated to have a relatively low homology with the known proteins in the same category. [0175] Herein, the term "relatively high homology" is defined as having 60% or more identity and the P-value 10<sup>-10</sup> or less in comparison with known sequences in the SwissProt database, or 64% or more identity and the P-value 10<sup>-15</sup> or less in comparison with those in the GenBank and UniGene databases (see the attached list). Also, the term "relatively low homology" is defined as not fulfilling the requirements to be "relatively high homology", but still having the scores, 25% or more identity and the P-value 10-6 or less, using the sequence having 55 nucleotides or more, in comparison with known sequences in the SwissProt database (see the attached list). The P-value is a score obtained statistically by taking into account the probability of occurrence of the similarity between two sequences. In general, the smaller P-value reflects the higher similarity (Altschul S.F., Gish W., Miller W., Myers E.W., and Lipman D.J. (1990) "Basic local alignment search tool" J.Mol. Biol., 215: 403-410; Gish W., and States D.J. (1993) "Identification of protein coding regions by database similarity search" Nature Genet. 3: 266-272).

[0176] The clones predicted to encode a protein in the category of secretory protein or membrane protein have the keywords, "signal", "transmembrane", "membrane", "extracellular matrix", "receptor", "G-protein coupled receptor", "ionic channel", "voltage-gated channel", "calcium channel", "cell adhesion", "collagen", or "connective tissue", or descriptions from which the clone can be predicted to be a secretory or membrane protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and

[0177] The clones predicted to encode a protein in the category of glycoprotein have the keywords, "glycoprotein", or descriptions from which the clone can be predicted to be a glycoprotein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence. [0178] The clones predicted to encode a protein in the category of proteins associated with signal transduction have the keywords, "senne/threonine-protein kinase", "tyrosine-protein kinase", "SH3 domain", or "WD repeat", or descriptions from which the clone can be predicted to be a protein associated with signal transduction (such as "ADP-ribosylation factor"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0179] The clones predicted to encode a protein in the category of proteins associated with transcription have the keywords, "transcription regulation", "zinc finger", or "homeobox", or descriptions from which the clone can be predicted to be a protein associated with transcription, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0180] The clones predicted to encode a protein in the category of proteins associated with diseases are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein that is deposited in the Online Mendelian Inheritance in Man (OMIM) database, which is a database of human genes and diseases, or the top hit data has descriptions from which the clone can be predicted to be a protein associated with diseases.

[0181] The clones predicted to encode a protein in the category of enzyme or proteins associated with metabolism are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein with E.C.No. (Enzyme commission number), or the top hit data has descriptions from which the clone can be predicted to be an enzyme or protein associated with metabolism (such as "metabolism", "oxidoreductase", or "lipid").

[0182] The clones predicted to encode a protein in the category of proteins associated with cell division or cell proliferation have the keywords, "cell division", "cell cycle", "mitosis", or "chromosomal protein", or descriptions from which the clone can be predicted to be a protein associated with cell division or cell proliferation (such as "histone", "cell growth", or "apoptosis"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0183] The clones predicted to encode a protein in the category of proteins associated with cytoskeleton have the keywords, "structural protein", "cytoskeleton", "actin-binding", or "microtubules", or descriptions from which the clone can be predicted to be a protein associated with cytoskeleton, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0184] The clones predicted to encode a protein in the category of proteins associated with RNA synthesis include the above clones predicted to be a protein associated with transcription, and also the clones which have the keywords, "RNA splicing", or "RNA processing", or descriptions from which the clone can be predicted to be a protein associated with RNA synthesis (such as "RNA helicase", "polyadenylation", or "RNA transport"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

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[0185] The clones predicted to encode a protein in the category of nuclear protein include the above clones predicted to be a protein associated with transcription, and also the clones which have the keyword, "nuclear protein", or descriptions from which the clone can be predicted to be a nuclear protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0186] The clones predicted to encode a protein in the category of proteins associated with protein synthesis or transport have the keywords, "translation regulation", "protein biosynthesis", "amino-acid biosynthesis", "ribosomal protein", or "protein transport", or descriptions from which the clone can be predicted to be a protein associated with protein synthesis or transport (such as "signal recognition particle", "ubiquitin", "proteosome", or "protease"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0187] The clones predicted to encode a protein in the category of proteins associated with cellular defense have the keywords, "heat shock", "chaperone", "DNA repair", or "DNA damage", or descriptions from which the clone can be predicted to be a protein associated with cellular defense, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0188] The clones predicted to encode a protein in the category of proteins associated with development or growth have the keyword, "developmental protein", or descriptions from which the clone can be predicted to be a protein associated with development or growth, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0189] The following 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth.

HEMBA1000055, HEMBA1000012, HEMBA1000020, HEMBA1000303, HEMBA1000158, HEMBA1000185, HEMBA1000201, HEMBA1000216, HEMBA1000303, HEMBA1000459, HEMBA1000488, HEMBA100049 1, HEMBA1000523, HEMBA1000531, HEMBA1000542, HEMBA1000561, HEMBA1000569, HEMBA10005 88, HEMBA1000852, HEMBA1000657, HEMBA1000673, HEMBA1000752, HEMBA1000827, HEMBA1000827, HEMBA1000851, HEMBA1001071, HEMBA1001071, HEMBA1001071, HEMBA1001071, HEMBA1001071, HEMBA1001073, HEMBA1001073, HEMBA1001073, HEMBA1001073, HEMBA1001073, HEMBA1001073, HEMBA1001074, HEMBA1001074, HEMBA1001074, HEMBA10010757, HEMBA1001369, HEMBA1001570, HEMBA1001570, HEMBA1001570, HEMBA1001595, HEMBA1001600, HEMBA1001600, HEMBA1001809, HEMBA1001819, HEMBA1001822, HEMBA1001847, HEMBA1001896, HEMBA1001913, HEMBA1002035, HEMBA1002035, HEMBA1002092, HEMBA1002022, HEMBA1002150, HEMBA1002341, HEMBA1002363, HEMBA1002417, HEMBA1002495, HEMBA1002513, HEMBA1002547, HEMBA1002555, HEMBA1002569, HEMBA1002609, HEMBA1002688, HEMBA1002716, HEMBA1002810, HEMBA1002896, HEMBA1002896,

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     [0191] The following 429 clones were estimated to have a relatively low homology with the known proteins in the
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same category.

	- HEMBA1000012, HEMBA1000752,	немва1000185,	HEMBA1000488,	HEMBA1000491,	HEMBA1000569,
5		HEMBA1001407, HEMBA1001809, HE		НЕМВА1001744,	HEMBA1001800,
10		HEMBA1002102, HEMBA1002609, HE			HEMBA1002495,
15				·	
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	HEMBA1003071, HEMBA1003291,	HEMBA1003077, HEMBA1003136, HEMBA1003179, HEMBA1003235, HEMBA1003418,
5	HEMBA1003568, HEMBA1003953,	HEMBA1003598, HEMBA1003680, HEMBA1003684, HEMBA1003836,
10	HEMBA1004276,	НЕМВА1004408, НЕМВА1004669, НЕМВА1004734,
	HEMBA1005475,	НЕМВА1005595, НЕМВА1005737, НЕМВА1005815, НЕМВА1005931,
15	HEMBA1006347, HEMBA1006737,	HEMBA1006474, HEMBA1006521, HEMBA1006562, HEMBA1006652, HEMBA1006914,
	HEMBA1007045,	НЕМВА1007121, НЕМВА1007300, НЕМВА1007301,
20	НЕМВВ1000119,	HEMBB1000217, HEMBB1000317, HEMBB1000632, HEMBB1000789,
	HEMBB1001011,	HEMBB1001177, HEMBB1001234, HEMBB1001282, HEMBB1001288,
25	HEMBB1002050,	HEMBB1002092,
	MAMMA1000085, MAMMA1001030,	MAMMA1000672, MAMMA1000841, MAMMA1000897, MAMMA1001008, MAMMA1001041,
30	MAMMA1002143, MAMMA1002671,	MAMMA1002573, MAMMA1002619, MAMMA1002636, MAMMA1002650, MAMMA1002881, MAMMA1002937,
	NT2RM1000118,	NT2RM1000153, NT2RM1000186, NT2RM1000187, NT2RM1000377,
35	NT2RM1000648, NT2RM1000850,	NT2RM1000666, NT2RM1000691, NT2RM1000770, NT2RM1000772, NT2RM1000882, NT2RM1000898,
40	NT2RM1001059,	NT2RM1001072, NT2RM1001139,
	NT2RM2000363, NT2RM2000623,	NT2RM2000371, NT2RM2000402, NT2RM2000577, NT2RM2000599, NT2RM2000714,
45	NT2RM2001131,	NT2RM2001221, NT2RM2001324, NT2RM2001345,
50	NT2RM2001588, NT2RM2001785, NT2RM2002580,	NT2RM2001635, NT2RM2001670, NT2RM2001700, NT2RM2001730, NT2RM2001797, NT2RM2001823, NT2RM2001936, NT2RM2001989,
. 55	NT2RM4000030, NT2RM4000191, NT2RM4000734,	NT2RM4000085, NT2RM4000155, NT2RM4000156, NT2RM4000169, NT2RM4000215, NT2RM4000354, NT2RM4000421, NT2RM4000496, NT2RM4000751,

_	NT2RM4001032, NT2RM4001444,	NT2RM4001140,	NT2RM4001316,	NT2RM4001340,	NT2RM4001410,
5	NT2RM4001611, NT2RM4001930,	NT2RM4001629, NT2RM4001979, NT2	NT2RM4001746, 2RM4001987,	NT2RM4001776,	NT2RM4001783,
10	NT2RM4002062,	NT2RM4002145, NT	2RM4002189, NT2R	M4002226, NT2RM4	002409,
	NT2RM4002527,	NT2RM4002571, NT	2RM4002623,		·
15	NT2RP1000111, NT2RP1000460,	NT2RP1000130, NT2RP1000470, NT	NT2RP1000333, 2RP1000493,	NT2RP1000348,	NT2RP1000363,
20	NT2RP1000547, NT2RP1001543,	NT2RP1000609,	NT2RP1001033,	NT2RP1001080,	NT2RP1001313,
20	NT2RP2000006, NT2RP2000448,	NT2RP2000133,	NT2RP2000161,	NT2RP2000183,	NT2RP2000257,
25	NT2RP2000710, NT2RP2001174,	NT2RP2000764, NT2RP2001839, NT	NT2RP2000812, 2RP2001900,	NT2RP2000932,	NT2RP2001127,
30	NT2RP2002025, NT2RP2002618,	NT2RP2002058, NT2RP2002710, NT	NT2RP2002185, 2RP2002980,	NT2RP2002442,	NT2RP2002595,
	NT2RP2003125, NT2RP2003564,	NT2RP2003277, NT2RP2003706,	NT2RP2003286,	NT2RP2003293,	NT2RP2003506,
35 40	NT2RP2004098, NT2RP2005037, NT2RP2005496, NT2RP2006275,	NT2RP2004187, NT2RP2005038, NT2RP2005531,	NT2RP2004568, NT2RP2005204, NT2RP2005690,	NT2RP2004791, NT2RP2005344, NT2RP2005763,	NT2RP2005022, NT2RP2005465, NT2RP2005767,
45	NT2RP3000046, NT2RP3000527, NT2RP3000994,	NT2RP3000047, NT2RP3000590,	NT2RP3000361, NT2RP3000603,	NT2RP3000397, NT2RP3000759,	NT2RP3000418, NT2RP3000868,
	NT2RP3001216, NT2RP3001426,	NT2RP3001274, NT2RP3001427,	NT2RP3001355,	NT2RP3001399,	NT2RP3001407,
50	NT2RP3001587, NT2RP3002399,	NT2RP3001708, NT2RP3002663, NT	NT2RP3002056, 2RP3002785, NT2F	NT2RP3002273, RP3002909,	NT2RP3002377,
55	NT2RP3003061, NT2RP3003825,	NT2RP3003500, NT2RP3003831,	NT2RP3003672,	NT2RP3003701,	NT2RP3003716,

	NT2RP3004209, NT2RP3004507,	NT2RP3004253, NT2RP3004569, NT2		NT2RP3004472, P3004617, NT2RP3	•
5	NT2RP4000210, NT2RP4000417, NT2RP4000879,	NT2RP4000212, NT2RP4000449, NT2RP4000925,	NT2RP4000259, NT2RP4000457,	NT2RP4000312, NT2RP4000498,	NT2RP4000370, NT2RP4000528,
10	NT2RP4001041, NT2RP4001149, NT2RP4001372,	NT2RP4001095, NT2RP4001150, NT2RP4001375,	NT2RP4001122, NT2RP4001174,	NT2RP4001126, NT2RP4001207,	NT2RP4001148, NT2RP4001219,
15	NT2RP4001551, NT2RP4001823, NT2RP4001946,	NT2RP4001592, NT2RP4001838,, NT2RP4002408,	NT2RP4001638, NT2RP4001849,	NT2RP4001725, NT2RP4001893,	NT2RP4001790, NT2RP4001927,
20	NT2RP5003477,	NT2RP5003522, NT2	RP5003524,		
25	OVARC1000004, OVARC1000746,	OVARC1000087, OVARC1000846, OVA	OVARC1000139, ARC1000876, OVAR	•	OVARC1000543, 000912,
	OVARC1001180, OVARC1001762,	OVARC1001329, OVARC1001880, OVA	OVARC1001476, ARC1001942, OVAR		•
30	PLACE1000066, PLACE1000863,	PLACE1000142,	PLACE1000420,	PLACE1000547,	PLACE1000583,
as	PLACE1001387,	PLACE1001632, PLA	CE1001692, PLAC	E1001989,	
35	PLACE1002450, PLACE1002991,	PLACE1002571,	PLACE1002722,	PLACE1002775,	PLACE1002811,
40	PLACE1003190, PLACE1003669,	PLACE1003258, PLACE1003704, PLA	PLACE1003420, CE1003915,	PLACE1003493,	PLACE1003605,
45	PLACE1004197, PLACE1004902,	PLACE1004428,	PLACE1004471,	PLACE1004629,	PLACE1004751,
	PLACE1005287,	PLACE1005557, PLA	CE1005611, PLACI	E1005763, PLACE10	005890,
50	PLACE1006157,	PLACE1006469, PLA	CE1006731,		
	PLACE1007238, PLACE1007729,	PLACE1007375, PLACE1008244, PLA	PLACE1007488, CE1008275, PLACE	PLACE1007544, E1008947,	PLACE1007598,
55	PLACE1009094,	PLACE1009319,	PLACE1009622,	PLACE1009721,	PLACE1009845,

PLACE1009861,

5	PLACE1010083, PLACE1010310,		PLACE1010106, PLACE1010960,	PLACE1010152, PLACE1011056,	PLACE1010194, PLACE1011219,
	PLACE1011263,	PLACE1011371, PLA	CE1011492,		
10	PLACE2000034, PLACE2000458,	PLACE2000164,	PLACE2000274,	PLACE2000399,	PLACE2000411,
	PLACE3000148,	PLACE3000156,	PLACE3000242,	PLACE3000322,	PLACE3000353,
15	PLACE3000373,				
	PLACE4000009.	PLACE4000014,	PLACE4000259,	PLACE4000431,	PLACE4000522,
		PLACE4000581, PLA		24000612,	•
20	1 MIOD 1000000,	. 2	,	·	
	SKNMC1000011,				
	THYR01000072.	THYR01000852, THY	/RO1000934, THYR	01000951,	
25	,			·	
	Y79AA1000258,	Y79AA1000368,	Y79AA1000539,	Y79AA1000589,	Y79AA1000962,
	Y79AA1001048,		Y79AA1001394,	Y79AA1001581,	Y79AA1001613,
	Y79AA1001692,		9AA1001874, Y79A	1001963,	
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Y79AA1002027, Y79AA1002139, Y79AA1002208, Y79AA1002209, Y79AA1002472, Y79AA1002482,

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(1) Among the 1430 clones, the following 259 clones were predicted to encode a protein in the category of secretory protein or membrane protein (including the clones belonging to plural categories). HEMBA1000005, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001351, HEMBA1001407, HEMBA1001476, HEMBA1001569, HEMBA1001678, HEMBA1001804, HEMBA1002150, HEMBA100254 7, HEMBA1002555, HEMBA1002688, HEMBA1002716, HEMBA1002999, HEMBA1003071, HEMBA10030 77, 40 HEMBA1003538, HEMBA1003598, HEMBA1003690, HEMBA1004207, HEMBA1004356, HEMBA1004 389, HEMBA1004756, HEMBA1005338, HEMBA1005367, HEMBA1005576, HEMBA1005699, HEMBA100 5963, HEMBA1005990, HEMBA1006310, HEMBA1006344, HEMBA1006562, HEMBA1006976, HEMBA10 07045. HEMBA1007301, HEMBB1000037, HEMBB1000119, HEMBB1000317, HEMBB1000530, HEMBB1 000915. HEMBB1001112, HEMBB1001119, HEMBB1001151, HEMBB1001282, HEMBB1001331, HEMBB 1002042. 45 HEMBB1002092, HEMBB1002193, HEMBB1002600, MAMMA1000103, MAMMA1000424, MAMM A1000672. MAMMA1000734, MAMMA1000897, MAMMA1001008, MAMMA1001030, MAMMA1001041, MAM MA1001075. MAMMA1001080, MAMMA1001751, MAMMA1001754, MAMMA1002143, MAMMA1002219, MA MMA1002485, MAMMA1002636, MAMMA1002938, MAMMA1003104, NT2RM1000199, NT2RM1000355, N T2RM1000430, 50 NT2RM1000648, NT2RM1000742, NT2RM1000770, NT2RM1000811, NT2RM1000833, NT2RM1000867, NT2RM1000882, NT2RM1000883, NT2RM1001139, NT2RM2000374, NT2RM2000402, NT2RM2000422, NT2RM2000566, NT2RM2000623, NT2RR2001324, NT2RM2001499, NT2RM200158 8, NT2RM2001592, NT2RM2001613, NT2RM2001635, NT2RM2001648, NT2RM2001760, NT2RM20021 09, NT2RM2002145. NT2RM4000156, NT2RM4000169, NT2RM4000386, NT2RM4000433, NT2RM4000795, NT2RM4001032. NT2RM4001054, NT2RM4001410, NT2RM4001605; NT2RM4001930, NT2RM4001987, NT2RM4002073, 55 NT2RM4002145, NT2RM4002189, NT2RM4002558, NT2RM400256 5, NT2RM4002571, NT2RM4002594, NT2RP1000130, NT2RP1000326, NT2RP1000363, NT2RP10004 13, NT2RP1000547, NT2RP1000782,

NT2RP1000856, NT2RP1001247, NT2RP1001313, NT2RP1001 546, NT2RP1001569, NT2RP2000056,

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NT2RP2000257, NT2RP2000842, NT2RP2001081, NT2RP200 1991, NT2RP2002025, NT2RP2002193,
         NT2RP2002256, NT2RP2002312, NT2RP2002479, NT2RP20 02504, NT2RP2003433, NT2RP2003706,
         NT2RP2003760, NT2RP2004194, NT2RP2004655, NT2RP2 005020, NT2RP2005344, NT2RP2005465,
         NT2RP2005752, NT2RP2005753, NT2RP2005835, NT2RP 2005933, NT2RP2006275, NT2RP2006565,
         NT2RP2006571, NT2RP3000324, NT2RP3000739, NT2R P3000919, NT2RP3001140, NT2RP3001355,
 5
         NT2RP3001676, NT2RP3001708, NT2RP3001727, NT2 RP3002659, NT2RP3003059, NT2RP3003101.
         NT2RP3003203, NT2RP3003212, NT2RP3003242, NT 2RP3003672, NT2RP3003701, NT2RP3003716,
         NT2RP3003918, NT2RP3004207, NT2RP3004253, N T2RP4000008, NT2RP4000212, NT2RP4000243,
         NT2RP4000417, NT2RP4000878, NT2RP4000907, NT2RP4000925, NT2RP4000928, NT2RP4001079,
         NT2RP4001117, NT2RP4001150, NT2RP4001174, NT2RP4001313, NT2RP4001315, NT2RP4001345,
1 10
         NT2RP4001372, NT2RP4001574, NT2RP400182 3, NT2RP4001966, NT2RP5003524, OVARC1000060,
         OVARC1000326, OVARC1000543, OVARC10006 82, OVARC1000722, OVARC1000912, OVARC1001055.
         OVARC1001154, OVARC1001261,
         OVARC1001329, OVARC1001419, OVARC1001506, OVARC1001542, OVARC1001610, OVARC1001713,
         OVARC1001802, OVARC1001943, PLACE1000611, PLACE1001092, PLACE1001136, PLACE100129 4.
 15
         PLACE1002437, PLACE1002500, PLACE1002722, PLACE1002782, PLACE1002816, PLACE10034 20,
         PLACE1003493, PLACE1003596, PLACE1004104, PLACE1004197, PLACE1004203, PLACE1004 277,
         PLACE1004629. PLACE1004751. PLACE1005494, PLACE1005804, PLACE1005890, PLACE100 6157,
         PLACE1007409. PLACE1007852. PLACE1008044. PLACE1008273. PLACE1008309. PLACE10 08603,
         PLACE1008643, PLACE1009659, PLACE1009670, PLACE1009845, PLACE1009861, PLACE1 011371,
 20
         PLACE1011492, PLACE1011896, PLACE2000034, PLACE2000341, PLACE2000399, PLACE 2000458,
         PLACE3000020, PLACE3000322, PLACE3000353, PLACE4000052, PLACE4000269, PLAC E4000522,
         PLACE4000581, THYRO1000783, Y79AA1000258, Y79AA1000346, Y79AA1001874,
         (2) The following 111 clones were predicted to encode a protein in the category of glycoprotein (including the clones
 25
         belonging to plural categories).
         HEMBA1000752, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001302, HEMBA1002150;
         HEMBA1002547, HEMBA1002555, HEMBA1003077, HEMBA1003538, HEMBA1003598, HEMBA100386 6,
         HEMBA1004356, HEMBA1005338, HEMBA1005576, HEMBA1005699, HEMBA1006562, HEMBA10069 76,
 30
         HEMBA1007301, HEMBB1000317, HEMBB1000530, HEMBB1001119, HEMBB1002092, HEMBB1002 193,
         MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA1001388, MAMMA1001771, MAMMA100 2485,
         MAMMA1002636, MAMMA1002938, NT2RM1000648, NT2RM2000374, NT2RM2000407, NT2RM20 00422,
         NT2RM2000623, NT2RM2001499, NT2RM2001588, NT2RM2001839, NT2RM2001930, NT2RM4 000156,
         NT2RM4000233, NT2RM4001410, NT2RM4001987, NT2RM4002145, NT2RM4002189, NT2RM 4002194,
         NT2RM4002558, NT2RM4002571, NT2RP1000856, NT2RP1001247, NT2RP2000056, NT2R P2001991,
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         NT2RP2002025, NT2RP2002325, NT2RP2002385, NT2RP2003706, NT2RP2005933, NT2 RP3001140,
         NT2RP3003242, NT2RP3003672, NT2RP3003701, NT2RP3003716, NT2RP3003914, NT 2RP3004253,
         NT2RP4000212, NT2RP4000417, NT2RP4000925, NT2RP4001149, NT2RP4001150, N T2RP4001219,
         NT2RP4001345, NT2RP4001372, NT2RP5003522, NT2RP5003524, OVARC1000543, OVARC1000682,
         OVARC1000722, OVARC1001055, OVARC1001154, OVARC1001506, OVARC1001713 PLACE1002437,
 40
         PLACE1002722, PLACE1003258, PLACE1003493, PLACE1004197, PLACE100475 1, PLACE1005698,
         PLACE1005804, PLACE1006157, PLACE1006754, PLACE1008643, PLACE10090 94, PLACE1009861,
         PLACE1011371, PLACE2000034, PLACE2000062, PLACE2000399, PLACE2000458, PLACE3000009,
         PLACE3000020, PLACE3000353, PLACE3000373, PLACE4000052, PLACE4000230, PLACE4000522,
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         PLACE4000581, THYRO1000852, Y79AA1001874,
         (3) The following 147 clones were predicted to encode a protein in the category of proteins associated with signal
         transduction (including the clones belonging to plural categories).
         HEMBA1000030, HEMBA1000185, HEMBA1000303, HEMBA1000459, HEMBA1000491, HEMBA1000657,
 50
         HEMBA1000991, HEMBA1001174, HEMBA1001387, HEMBA1001822, HEMBA1001921, HEMBA100221 2,
         HEMBA1002341, HEMBA1002417, HEMBA1002810, HEMBA1002896, HEMBA1003291, HEMBA10033 14,
         HEMBA1003560, HEMBA1003615, HEMBA1003621, HEMBA1003805, HEMBA1004286, HEMBA1004 604,
         HEMBA1006173, HEMBA1006540, HEMBA1006973, HEMBA1007174, HEMBB1000632, HEMBB100 0781,
         HEMBB1001051, HEMBB1001177, HEMBB1001234, HEMBB1001294, HEMBB1001384, HEMBB10 02193,
 55
         HEMBB1002477, HEMBB1002635, MAMMA1001305, MAMMA1002699, MAMMA1002842, MAMMA1 003057,
         MAMMA1003113, NT2RM1000118, NT2RM1000186, NT2RM1000772, NT2RM1000850, NT2RM 1001072,
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NT2RM2000124, NT2RM2000612, NT2RM2001065, NT2RM2001221, NT2RM2001345, NT2R M2001652, NT2RM2001983, NT2RM2002109, NT2RM2002128, NT2RM4000327, NT2RM4000354, NT2 RM4000798.

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NT2RM4001203, NT2RM4001411, NT2RM4001412, NT2RM4001582, NT2RM4001629, NT 2RM4001758.
       NT2RM4002226. NT2RP1000018, NT2RP1000112, NT2RP1000701, NT2RP1000825, N T2RP1001395,
       NT2RP1001410. NT2RP1001457, NT2RP1001482, NT2RP2002058, NT2RP2002710, NT2RP2003702,
       NT2RP2003912, NT2RP2004232, NT2RP2004768, NT2RP2005022, NT2RP2005620, NT2RP2005890,
       NT2RP3000299, NT2RP3000403, NT2RP3000742, NT2RP3000845, NT2RP300176 4, NT2RP3002004,
       NT2RP3002785, NT2RP3002909, NT2RP3003230, NT2RP3003411, NT2RP30034 64, NT2RP3003800.
       NT2RP3004475, NT2RP3004534, NT2RP4000147, NT2RP4000376,
       NT2RP4001122, NT2RP4001375, NT2RP4001644, NT2RP4001725, NT2RP4001760, NT2RP4001849,
       NT2RP4001927, NT2RP4002408, NT2RP5003477, OVARC1000437, OVARC1000465, OVARC100064 9,
       OVARC1000945, OVARC1001476, OVARC1001861, PLACE1001377, PLACE1001384, PLACE10013 87,
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       PLACE1002493, PLACE1002591, PLACE1002685, PLACE1003353, PLACE1004128, PLACE1004 918,
       PLACE1005502, PLACE1005739, PLACE1005968, PLACE1006385, PLACE1007238, PLACE100 7375,
       PLACE1007488, PLACE1008244, PLACE1008650, PLACE1009319, PLACE1009468, PLACE10 10083,
       PLACE1010942, PLACE1011041, PLACE1011046, PLACE1011923, PLACE2000164, PLACE3 000145,
       PLACE3000350, PLACE3000475, THYRO1001595, Y79AA1000059, Y79AA1000966,
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(4) The following 230 clones were predicted to encode a protein in the category of proteins associated with transcription (including the clones belonging to plural categories). HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000851, HEMBA1001137, HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA100184 7, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA1003545, HEMBA10035 68, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004 353, HEMBA1004479, HEMBA1004758, HEMBA1005359, HEMBA1005528, HEMBA1005548, HEMBA100 5931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006559, HEMBB1000250, HEMBB10 00789, HEMBB1001011, HEMBB1001314, HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1 002217. MAMMA1000155, MAMMA1000183, MAMMA1000388, MAMMA1001633, MAMMA1001820, MAMMA 1001837, MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000666, NT2RM1000691, NT2R M1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2 RM2001605. NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2002580, NT2RM4000024, NT 2RM4000104, NT2RM4000191, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, N T2RM4000996,  $NT2RM4001140, \quad NT2RM4001200, \quad NT2RM4001483, \quad NT2RM4001592, \quad NT2RM4001746, \quad NT2RM4001783, \quad NT2RM4001783$ NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RP1000111, NT2RP1000333, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP2000008, NT2RP200013 3, NT2RP2000297. NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2002252, NT2RP20025 03, NT2RP2002591, NT2RP2003125, NT2RP2003228, NT2RP2003293, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004187, NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP2006238, NT2RP200646 4, NT2RP3000050, NT2RP3000397, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP30006 32, NT2RP3001057, NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001 399, NT2RP3001671, NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP300 1855, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP30 03251, NT2RP3003327, NT2RP3004258, NT2RP3004566, NT2RP3004572, NT2RP3004594, NT2RP3 004617, NT2RP4000078, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000787, NT2RP 4000865, NT2RP4000997, NT2RP4001029, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2R P4001551, NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVA RC1000479, OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577, OVARC1001949, PL ACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706, PLACE1001118, P LACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE1002450, PLACE1002532, PLACE1002834, PLACE1003302, PLACE1003605, PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004681, PLACE1006414, PLACE1006438, PLACE1006482, PLACE100723 9, PLACE1007346, PLACE1007544, PLACE1007598, PLACE1007955, PLACE1007969, PLACE10086 27, PLACE1008941,

PLACE1009099, PLACE1009246, PLACE1009398, PLACE1010261,
PLACE1010491, PLACE1010702, PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072,
PLACE2000359, PLACE4000128, PLACE4000156, PLACE4000259, PLACE4000431, THYRO100012 1,
THYRO1000242, THYRO1000501, THYRO1000585, THYRO1001100, THYRO1001189, Y79AA10000 37,
Y79AA1000627, Y79AA1000705, Y79AA1001105, Y79AA1001299, Y79AA1001391, Y79AA1001 533,
Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1002093, Y79AA1002472, Y79AA100 2482,

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(5) The following 436 clones were predicted to encode a protein in the category of proteins associated with diseases
       (including the clones belonging to plural categories).
       HEMBA1000158, HEMBA1000216, HEMBA1000523, HEMBA1000561, HEMBA1000569, HEMBA1000673,
       HEMBA1000972, HEMBA1001017, HEMBA1001019, HEMBA1001059, HEMBA1001071, HEMBA100108 8,
       HEMBA1001123, HEMBA1001407, HEMBA1001569, HEMBA1001570, HEMBA1001672, HEMBA10020 35,
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       HEMBA1002160, HEMBA1002162, HEMBA1002547, HEMBA1002939, HEMBA1003077, HEMBA1003 418,
       HEMBA1003433, HEMBA1003538, HEMBA1003555, HEMBA1003568, HEMBA1003569, HEMBA100 3581,
       HEMBA1003720, HEMBA1003760, HEMBA1004168, HEMBA1004248, HEMBA1004276, HEMBA10 04354,
       HEMBA1004356, HEMBA1004479, HEMBA1004534, HEMBA1004669, HEMBA1004752, HEMBA1 004753.
       HEMBA1004807, HEMBA1005293, HEMBA1005331, HEMBA1005359, HEMBA1005367, HEMBA 1005423,
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       HEMBA1005443, HEMBA1005475, HEMBA1005528, HEMBA1005576, HEMBA1005581, HEMB A1005679,
       HEMBA1005699, HEMBA1005732, HEMBA1005931, HEMBA1006173, HEMBA1006474, HEM BA1006562,
       HEMBA1006648, HEMBA1006652, HEMBA1006737, HEMBA1006807, HEMBA1006976, HE MBA1007078,
       HEMBA1007121, HEMBA1007243, HEMBA1007301, HEMBB1000264, HEMBB1000637, H EMBB1000684.
       HEMBB1000693, HEMBB1000915, HEMBB1000927, HEMBB1001011, HEMBB1001051, HEMBB1001068,
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       HEMBB1001102, HEMBB1001119, HEMBB1001133, HEMBB1001177, HEMBB1001802, HEMBB1001908,
       HEMBB1001922, HEMBB1001967, HEMBB1002050, HEMBB1002193, HEMBB100221 7, HEMBB1002358,
       HEMBB1002635, HEMBB1002683, MAMMA100009, MAMMA100092, MAMMA10001 03, MAMMA1000173,
       MAMMA1000731, MAMMA1000897, MAMMA1000956, MAMMA1001021,
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        MAMMA1001041, MAMMA1001105, MAMMA1001198, MAMMA1001305, MAMMA1001576, MAMMA1001633,
        MAMMA1002140, MAMMA1002170, MAMMA1002198, MAMMA1002268, MAMMA1002352, MAMMA100239 2,
        MAMMA1002485, MAMMA1002530, MAMMA1002573, MAMMA1002636, MAMMA1002858, MAMMA10028 69.
        MAMMA1002881, MAMMA1002937, MAMMA1003047, MAMMA1003099, MAMMA1003146, MAMMA1003 166,
        NT2RM1000035, NT2RM1000153, NT2RM1000257, NT2RM1000318, NT2RM1000377, NT2RM100 0394,
        NT2RM1000555, NT2RM1000691, NT2RM1000725, NT2RM1000770, NT2RM1000826, NT2RM10 00850,
25
        NT2RM1001059, NT2RM1001072, NT2RM1001082, NT2RM1001092, NT2RM2000363, NT2RM2 000374,
        NT2RM2000594, NT2RM2000599, NT2RM2000714, NT2RM2000740, NT2RM2000821, NT2RM 2001035,
        NT2RM2001499, NT2RM2001575, NT2RM2001605, NT2RM2001664, NT2RM2001670, NT2R M2001803,
        NT2RM2001823, NT2RM2001839, NT2RM4000085, NT2RM4000139, NT2RM4000155, NT2 RM4000290,
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        NT2RM4000689, NT2RM4000895, NT2RM4001320, NT2RM4001566, NT2RM4001629, NT 2RM4001741,
        NT2RM4001819, NT2RM4001823, NT2RM4001865, NT2RM4001940, NT2RM4002093, N T2RM4002146,
        NT2RM4002161, NT2RM4002189, NT2RM4002457, NT2RM4002558, NT2RM4002571, NT2RP1000086,
        NT2RP1000376, NT2RP1000574, NT2RP1000581, NT2RP1000629, NT2RP1000721, NT2RP1000733,
        NT2RP1000738, NT2RP1000825, NT2RP1000833, NT2RP1000860, NT2RP100096 6, NT2RP1001033,
        NT2RP1001247, NT2RP2000126, NT2RP2000147, NT2RP2000233, NT2RP20002 97, NT2RP2000310,
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        NT2RP2000414, NT2RP2000448, NT2RP2000459, NT2RP2000603,
        NT2RP2000812, NT2RP2001127, NT2RP2001327, NT2RP2001440, NT2RP2001520, NT2RP2001663,
        NT2RP2001748, NT2RP2001876, NT2RP2001898, NT2RP2002025, NT2RP2002066, NT2RP200225 9,
        NT2RP2002312, NT2RP2002503, NT2RP2002618, NT2RP2003228, NT2RP2003293, NT2RP20032 95,
        NT2RP2003517, NT2RP2003564, NT2RP2003706, NT2RP2003737, NT2RP2003968, NT2RP2004 013,
40
        NT2RP2004098, NT2RP2004187, NT2RP2004316, NT2RP2004523, NT2RP2004538, NT2RP200 4933,
        NT2RP2005144, NT2RP2005288, NT2RP2005289, NT2RP2005325, NT2RP2005358, NT2RP20 05531,
        NT2RP2005773, NT2RP2006219, NT2RP3000050, NT2RP3000080, NT2RP3000299, NT2RP3 000397,
        NT2RP3000512, NT2RP3000527, NT2RP3001081, NT2RP3001216, NT2RP3001268, NT2RP 3001355,
        NT2RP3001427, NT2RP3001428, NT2RP3001497, NT2RP3001554, NT2RP3001724, NT2R P3001764,
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        NT2RP3001799, NT2RP3001855, NT2RP3002045, NT2RP3002151, NT2RP3002351, NT2 RP3002352,
        NT2RP3002663, NT2RP3002818, NT2RP3002876, NT2RP3002909, NT2RP3002972, NT 2RP3003061,
        NT2RP3003157, NT2RP3003242, NT2RP3003282, NT2RP3003672, NT2RP3004078, N T2RP3004209,
        NT2RP3004399, NT2RP3004424, NT2RP3004475, NT2RP3004480, NT2RP3004490, NT2RP3004527.
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        NT2RP3004578, NT2RP4000049, NT2RP4000109, NT2RP4000185, NT2RP4000312, NT2RP4000367,
        NT2RP4000398, NT2RP4000457, NT2RP4000855, NT2RP4000879, NT2RP400091 5, NT2RP4000928,
        NT2RP4001095, NT2RP4001150, NT2RP4001213, NT2RP4001483, NT2RP40014 98, NT2RP4001753,
        NT2RP4001838, OVARC1000014, OVARC1000114, OVARC1000139,
        OVARC1000241, OVARC1000440, OVARC1000543, OVARC1000722, OVARC1000771, OVARC1000834,
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        OVARC1000850, OVARC1001113, OVARC1001244, OVARC1001296, OVARC1001357, OVARC100137 2,
        OVARC1001417, OVARC1001496, OVARC1001506, OVARC1001577, OVARC1001668, OVARC10017 13,
        OVARC1001809. OVARC1001880. OVARC1001901. OVARC1001949. OVARC1002050. PLACE1000 133.
        PLACE1000142. PLACE1000184. PLACE1000383. PLACE1000406. PLACE1000420. PLACE100 0583.
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PLACE1000588, PLACE1000706, PLACE1001384, PLACE1001387, PLACE1001517, PLACE10 01602.
       PLACE1001634, PLACE1001771, PLACE1002046, PLACE1002437, PLACE1002474, PLACE1 002665,
       PLACE1002834, PLACE1003302, PLACE1003493, PLACE1003704, PLACE1003888, PLACE 1003903,
       PLACE1003968, PLACE1004197, PLACE1004358, PLACE1004471, PLACE1004681, PLAC E1004751,
       PLACE1004773, PLACE1005101, PLACE1006040, PLACE1006119, PLACE1006170, PLA CE1006187.
       PLACE1006288, PLACE1006414, PLACE1006438, PLACE1006962, PLACE1007239, PL ACE1007257,
       PLACE1007488, PLACE1007511, PLACE1007737, PLACE1008000, PLACE1008402, P LACE1008643,
       PLACE1008696, PLACE1008941, PLACE1009027, PLACE1009200, PLACE1009230, PLACE1009298,
       PLACE1009319, PLACE1009444, PLACE1009524, PLACE1010083, PLACE1010089, PLACE1010599,
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       PLACE1010833, PLACE1010856, PLACE1011054, PLACE1011057, PLACE101116 5, PLACE1011229,
       PLACE1011263, PLACE1011371, PLACE2000072, PLACE2000399, PLACE20004 38, PLACE2000458,
       PLACE3000004, PLACE3000242, PLACE3000352, PLACE3000353,
       PLACE3000455, PLACE3000475, PLACE4000009, PLACE4000014, PLACE4000052, PLACE4000320,
       PLACE4000581, SKNMC1000050, THYRO1000026, THYRO1000173, THYRO1000186, THYRO100032 0,
15
       THYRO1000327, THYRO1000387, THYRO1000471, THYRO1000934, THYRO1001458, THYRO10015 95,
       THYRO1001605, THYRO1001617, THYRO1001671, Y79AA1000037, Y79AA1000469, Y79AA1000 539,
       Y79AA1000560, Y79AA1000734, Y79AA1000985, Y79AA1001105, Y79AA1001391, Y79AA100 1394,
       Y79AA1001548, Y79AA1001613, Y79AA1001874, Y79AA1002258, Y79AA1002416, Y79AA10 02482,
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20 (6) The following 301 clones were predicted to encode a protein in the category of enzyme or proteins associated with metabolism (including the clones belonging to plural categories). HEMBA1000012, HEMBA1000459, HEMBA1000852, HEMBA1001059, HEMBA1001257, HEMBA1001620, HEMBA1001714, HEMBA1001744, HEMBA1001896, HEMBA1002003, HEMBA1002212, HEMBA100225 7, HEMBA1002513, HEMBA1002973, HEMBA1003046, HEMBA1003136, HEMBA1003179, HEMBA10032 86, 25 HEMBA1003291, HEMBA1003538, HEMBA1003680, HEMBA1004227, HEMBA1004408, HEMBA1004 734, HEMBA1004795, HEMBA1005513, HEMBA1005737, HEMBA1005815, HEMBA1006278, HEMBA100 6347, HEMBA1006521, HEMBA1006648, HEMBA1006877, HEMBA1006941, HEMBA1006976, HEMBA10 07121, HEMBA1007243, HEMBA1007300, HEMBB1000781, HEMBB1000915, HEMBB1001288, HEMBB1 001429, HEMBB1001443, HEMBB1001915, HEMBB1002042, HEMBB1002342, HEMBB1002358, HEMBB 1002635, 30 HEMBB1002683, MAMMA1000020, MAMMA1000085, MAMMA1000672, MAMMA1000841, MAMM A1001008, MAMMA1001139, MAMMA1001476, MAMMA1001501, MAMMA1002268, MAMMA1002530, MAM MA1002619, MAMMA1002671, MAMMA1002938, NT2RM1000132, NT2RM1000153, NT2RM1000256, NT 2RM1000280, NT2RM1000377, NT2RM1000648, NT2RM1000850, NT2RM1000867, NT2RM1000894, N T2RM1001072, NT2RM2000013, NT2RM2000124, NT2RM2000191, NT2RM2000368, NT2RM2000371, NT2RM2000577, NT2RM2000594, NT2RM2000599, NT2RM2000951, NT2RM2001238, NT2RM2001664, NT2RM2001700, 35 NT2RM2001730, NT2RM2001785, NT2RM2001803, NT2RM2002030, NT2RM400002 4, NT2RM4000155, NT2RM4000344, NT2RM4000496, NT2RM4000616, NT2RM4000712, NT2RM40008 95, NT2RM4001313, NT2RM4001316, NT2RM4001340, NT2RM4001444, NT2RM4001758, NT2RM4001819, NT2RM4001930, NT2RM4002062, NT2RM4002063, NT2RM4002409, NT2RM4002558, 40 NT2RM4002571, NT2RM4002623, NT2RP1000018, NT2RP1000112, NT2RP1000376, NT2RP100060 9, NT2RP1000833, NT2RP1000834, NT2RP1001079, NT2RP1001253, NT2RP1001361, NT2RP10015 43, NT2RP2000056, NT2RP2000114, NT2RP2000183, NT2RP2000329, NT2RP2000422, NT2RP2000 710, NT2RP2000764, NT2RP2000932, NT2RP2001070, NT2RP2001663, NT2RP2001839, NT2RP200 1898, NT2RP2002256, NT2RP2002312, NT2RP2002442, NT2RP2002595, NT2RP2002618, NT2RP20 02993, 45 NT2RP2003230, NT2RP2003286, NT2RP2003506, NT2RP2003643, NT2RP2003706, NT2RP2 003737, NT2RP2003781, NT2RP2003912, NT2RP2004232, NT2RP2004239, NT2RP2004768, NT2RP 2004791, NT2RP2004799, NT2RP2005038, NT2RP2005276, NT2RP2005344, NT2RP2005457, NT2R P2005498, NT2RP2005531, NT2RP2005557, NT2RP2005690, NT2RP2005773, NT2RP2005775, NT2 RP2005942, NT2RP2006571, NT2RP3000046, NT2RP3000085, NT2RP3000359, NT2RP3000418, NT 2RP3000742, 50 NT2RP3000845, NT2RP3000875, NT2RP3001274, NT2RP3001407, NT2RP3001495, N T2RP3002351, NT2RP3002402, NT2RP3002969, NT2RP3003301, NT2RP3003346, NT2RP3003500, NT2RP3003800, NT2RP3003825, NT2RP3003831, NT2RP3003914, NT2RP3004155, NT2RP3004209 NT2RP3004669, NT2RP3004670, NT2RP4000259, NT2RP4000417, NT2RP4000457, NT2RP400072 4, NT2RP4000855, NT2RP4000928, NT2RP4000997, NT2RP4001041, NT2RP4001079, NT2RP40010 95, NT2RP4001219, 55 NT2RP4001345, NT2RP4001375, NT2RP4001483, NT2RP4001592, NT2RP4001644, NT2RP4001893, NT2RP4001946, NT2RP4002408, NT2RP5003500, NT2RP5003522, OVARC1000060, OVARC1000139, OVARC1000309, OVARC1000543, OVARC1000682, OVARC100072 2,

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OVARC1001942, PLACE1000007, PLACE1000142, PLACE1000420, PLACE1000547, PLACE1000 653, PLACE1001062, PLACE1001603, PLACE1001692, PLACE1001817, PLACE1001989, PLACE100 2991, PLACE1003908, PLACE1003709, PLACE1003885, PLACE1003888, PLACE1003903, PLACE10 03915, PLACE1005656, PLACE1005763, PLACE1005804, PLACE1004437, PLACE1005656, PLACE1005763, PLACE1005804, PLACE1006011, PLACE1016469, PLACE 1006731, PLACE1007729, PLACE1007958, PLACE1008275, PLACE1008356, PLACE1008696, PLAC E1009444, PLACE1010481, PLACE1011046, PLACE10110089, PLACE10110096, PLACE1011006, PLA CE101152, PLACE1011923, PLACE1011046, PLACE1011203, PLACE1011219, PLACE1011229, PLACE1011332, PLACE3000148, PLACE3000156, PLACE3000350, PLACE3000353, PLACE4000259, PLACE4000431, PLACE4000558, THYRO1000358, THYRO1000471, THYRO1000926, THYRO1000934, THYRO100095 1, THYRO1000320, THYRO1001617, THYRO1000671, Y79AA1001048, Y79AA1001233, Y79AA10015 81, Y79AA1001679, Y79AA1001827, Y79AA1002027, Y79AA1002209, Y79AA1002361, Y79AA1002 416,
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- (7) The following 74 clones were predicted to encode a protein in the category of proteins associated with cell division or cell proliferation (including the clones belonging to plural categories).

  HEMBA1001019, HEMBA1001595, HEMBA1002229, HEMBA1002363, HEMBA1003136, HEMBA1003836, HEMBA1004131, HEMBA1005241, HEMBB1000399, HEMBB1001175, HEMBB1001242, MAMMA100182 0, MAMMA1003011, NT2RM1000394, NT2RM2001256, NT2RM4000215, NT2RM4001611, NT2RM40017 14, NT2RP1000733, NT2RP1001177, NT2RP2000077, NT2RP2000346, NT2RP2001397, NT2RP2003 912, NT2RP2005520, NT2RP2005675, NT2RP2005767, NT2RP2005857, NT2RP2006464, NT2RP300 0031, NT2RP3001155, NT2RP3001730, NT2RP3002151, NT2RP3002273, NT2RP3002399, NT2RP30 04507, NT2RP3004578, NT2RP3004594, NT2RP4000210, NT2RP4000498, NT2RP4001207, NT2RP4 001414, NT2RP4001551, OVARC1000066, OVARC1000087, OVARC1000113, OVARC1000846, OVARC 1000876, OVARC1001068, OVARC1001107, OVARC1001113, OVARC1002112, PLACE100547, PLAC E1001611, PLACE1004814, PLACE1005287, PLACE1006187, PLACE1003709, PLACE1004316, PLA CE1004674, PLACE100960, PLACE1010720, PLACE1010761, PLACE1010833, PLACE1011056, P LACE3000244, Y79AA1000214, Y79AA1000214, Y79AA1001236,
- (8) The following 92 clones were predicted to encode a protein in the category of proteins associated with cytoskeleton (including the clones belonging to plural categories).
- HEMBA1000020, HEMBA1000588, HEMBA1001804, HEMBA1002102, HEMBA1002161, HEMBA1002939. HEMBA1003235, HEMBA1003581, HEMBA1004534, HEMBA1005009, HEMBA1005595, HEMBA100627 2, HEMBA1006737, HEMBA1007018, HEMBB1000404, HEMBB1001282, MAMMA1000173, MAMMA10004 29, MAMMA1001041, MAMMA1001576, MAMMA1001735, MAMMA1002622, MAMMA1002637, MAMMA1003 099, MAMMA1003127, NT2RM1000898, NT2RM1001003, NT2RM1001139, NT2RM2000691, NT2RM400 0030, NT2RM4000167, NT2RM4000689, NT2RM4001741, NT2RM4001776, NT2RM4002109, NT2RP10 00348, NT2RP1000460, NT2RP1000478, NT2RP1001033, NT2RP2000812, NT2RP2001634, NT2RP2 001900. NT2RP2002727, NT2RP2003307, NT2RP2003604, NT2RP2004538, NT2RP2005531, NT2RP 3000868, NT2RP3001216, NT2RP3001272, NT2RP3002688, NT2RP3003061, NT2RP3003138, NT2R P3003230, NT2RP3003282, NT2RP3004454, NT2RP3004569, NT2RP4001126, NT2RP4001927, OVA RC1000437, OVARC1000520, OVARC1000679, OVARC1001731, OVARC1002050, PLACE1002571, PL ACE1002591, PLACE1002655, PLACE1003669, PLACE1004777, PLACE1005287, PLACE1006368, P LACE1007105, PLACE1010310, PLACE1010743, PLACE1010960, PLACE1011263, PLACE2000039, PLACE2000216, PLACE2000274, PLACE3000145, PLACE3000322, PLACE4000009, PLACE4000612, THYRO1000072, THYRO1000666, THYRO1001121, THYRO1001458, Y79AA1000368, Y79AA100083 3, Y79AA1000962, Y79AA1000985, Y79AA1002208,

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- (9) The following 280 clones were predicted to encode a protein in the category of proteins associated with RNA synthesis (including the clones belonging to plural categories).
- HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561. HEMBA1000591, HEMBA1000851, HEMBA1001137, HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA100181 9, HEMBA1001847, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA10034 18, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004 097, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA1004669, HEMBA1004758, HEMBA100 5359, HEMBA1005528, HEMBA1005548, HEMBA1005931, HEMBA1006253, HEMBA1006278, HEMBA10 06359,

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HEMBA1006559, HEMBA1006639, HEMBB1000250, HEMBB1000789, HEMBB1001011, HEMBB1 001314,
       HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAMMA 1000183,
       MAMMA1000388, MAMMA1001059, MAMMA1001633, MAMMA1001820, MAMMA1001837, MAMM A1002650,
       MAMMA1002937, NT2RM1000055, NT2RM1000187, NT2RM1000666, NT2RM1000691, NT2 RM1000852,
       NT2RM1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT 2RM2001424,
       NT2RM2001575, NT2RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001797, N T2RM2001989,
       NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000104, NT2RM4000191, NT2RM4000421,
       NT2RM4000471, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, NT2RM4000996,
       NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM400174 6, NT2RM4001783,
       NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RM40020 93, NT2RM4002479,
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       NT2RP1000111, NT2RP1000272, NT2RP1000333, NT2RP1000470,
       NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1001080, NT2RP2000008, NT2RP2000133.
       NT2RP2000161, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP200144 9,
       NT2RP2002099, NT2RP2002252, NT2RP2002503, NT2RP2002591, NT2RP2002928, NT2RP20031 25,
       NT2RP2003228, NT2RP2003293, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004 187,
15
       NT2RP2004568, NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005126, NT2RP200 5168,
       NT2RP2005239, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP20 05763,
       NT2RP2005942, NT2RP2006238, NT2RP2006464, NT2RP3000050, NT2RP3000361, NT2RP3 000397,
       NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP3000632, NT2RP3000917, NT2RP 3001057,
       NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001399, NT2R P3001671,
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       NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP3001855, NT2 RP3002165,
       NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP3003251, NT 2RP3003327,
       NT2RP3004013, NT2RP3004258, NT2RP3004504, NT2RP3004566, NT2RP3004572, N T2RP3004594,
       NT2RP3004617, NT2RP4000078, NT2RP4000111, NT2RP4000210, NT2RP4000398, NT2RP4000449,
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       NT2RP4000518, NT2RP4000614, NT2RP4000787, NT2RP4000865, NT2RP4000997, NT2RP4001029,
       NT2RP4001080, NT2RP4001148, NT2RP4001213, NT2RP4001433, NT2RP400152 9, NT2RP4001551,
       NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP40020 78, OVARC1000479,
       OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577,
       OVARC1001949, PLACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706,
       PLACE1001118, PLACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE100245 0,
30
       PLACE1002532, PLACE1002834, PLACE1003190, PLACE1003302, PLACE1003605, PLACE10037 04,
       PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004 681,
       PLACE1004902, PLACE1005646, PLACE1005876, PLACE1006196, PLACE1006414, PLACE100 6438,
       PLACE1006482, PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10 07955,
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       PLACE1007969, PLACE1008627, PLACE1008941, PLACE1009099, PLACE1009246, PLACE1 009398,
       PLACE1010053, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010580, PLACE 1010702,
       PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072, PLACE2000359, PLAC E4000128,
       PLACE4000156, PLACE4000259, PLACE4000431, THYRO1000121, THYRO1000242, THY RO1000501,
       THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000349, Y7 9AA1000539,
       Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1001105, Y79AA1001299, Y 79AA1001391,
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       Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1001963, Y79AA1002093,
       Y79AA1002472, Y79AA1002482,
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(10) The following 352 clones were predicted to encode a protein in the category of nuclear protein (including the clones belonging to plural categories).

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HEMBA100005, HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000827, HEMBA1000851, HEMBA1001088, HEMBA1001137, HEMBA1001476, HEMBA1001510, HEMBA100157 9, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1002035, HEMBA10020 92, HEMBA1002363, HEMBA1002495, HEMBA1002569, HEMBA1002609, HEMBA1002999, HEMBA1003 148, HEMBA1003418, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA100 3783, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA10 04669, HEMBA1004758, HEMBA1005359, HEMBA1005475, HEMBA1005528, HEMBA1005548, HEMBA1 005931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006474, HEMBA1006559, HEMBA 1006914, HEMBB1001908, HEMBB1000399, HEMBB1000789, HEMBB1001011, HEMBB1001314, HEMB B1001482, HEMBB1001908, HEMBB1001915, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAM MA1000183, MAMMA1000388, MAMMA1000731, MAMMA1001633, MAMMA1001820, MAMMA1001837, N T2RM1000257, NT2RM1000394, NT2RM1000666, NT2RM1000691, NT2RM1000852, NT2RM1000894, NT2RM1001092,

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NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2RM2001605, NT2RM2001635,
       NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2001823, NT2RM200193 6, NT2RM2001989,
       NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000085, NT2RM40001 04, NT2RM4000191,
       NT2RM4000215, NT2RM4000290, NT2RM4000421, NT2RM4000496,
       NT2RM4000531, NT2RM4000734, NT2RM4000751, NT2RM4000996, NT2RM4001140, NT2RM4001200,
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       NT2RM4001217, NT2RM4001382, NT2RM4001483, NT2RM4001592, NT2RM4001605, NT2RM400161 1,
       NT2RM4001746, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM40019 79.
       NT2RM4002093, NT2RM4002146, NT2RM4002527, NT2RP1000035, NT2RP1000111, NT2RP1000 333,
       NT2RP1000493, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1000966, NT2RP100 1177,
       NT2RP2000008, NT2RP2000126, NT2RP2000133, NT2RP2000161, NT2RP2000297, NT2RP20 00414,
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       NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2001420, NT2RP2001449, NT2RP2 001536,
       NT2RP2001762, NT2RP2002185, NT2RP2002252, NT2RP2002259, NT2RP2002503, NT2RP 2002504,
        NT2RP2002591, NT2RP2002993, NT2RP2003125, NT2RP2003228, NT2RP2003277, NT2R P2003286,
       NT2RP2003293, NT2RP2003308, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2 RP2004187,
       NT2RP2004961,\ NT2RP2005003,\ NT2RP2005037,\ NT2RP2005287,\ NT2RP2005496,\ NT\ 2RP2005520,
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       NT2RP2005539, NT2RP2005722, NT2RP2005767, NT2RP2005857, NT2RP2005942, N T2RP2006238.
        NT2RP3006464, NT2RP3000031, NT2RP3000050, NT2RP3000361, NT2RP3000397, NT2RP3000512.
        NT2RP3000527, NT2RP1000590, NT2RP3000603, NT2RP3000632, NT2RP3000917, NT2RP3001057,
        NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001274, NT2RP300139 8, NT2RP3001399,
        NT2RP3001427, NT2RP3001428, NT2RP3001587, NT2RP3001671, NT2RP30016 72, NT2RP3001688,
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        NT2RP3001712, NT2RP3001724, NT2RP3001792, NT2RP3001855,
        NT2RP3002056, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193,
        NT2RP3003212, NT2RP3003251, NT2RP3003327, NT2RP3004206, NT2RP3004209, NT2RP300425 8,
        NT2RP3004566, NT2RP3004572, NT2RP3004594, NT2RP3004617, NT2RP4000078, NT2RP40001 11,
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        NT2RP4000150, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000457, NT2RP4000 518,
        NT2RP4000787, NT2RP4000865, NT2RP4000997, NT2RP4001029, NT2RP4001080, NT2RP400 1148,
        NT2RP4001207, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2RP4001551, NT2RP40 01638,
        NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVARC1000006, OVARC1 000087,
        OVARC1000139, OVARC1000326, OVARC1000440, OVARC1000479, OVARC1000846, OVARC 1000883,
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        OVARC1001180, OVARC1001244, OVARC1001381, OVARC1001555, OVARC1001577, OVAR C1001702,
        OVARC1001949, OVARC1002112, PLACE1000066, PLACE1000133, PLACE1000406, PLA CE1000583,
        PLACE1000596, PLACE1000706, PLACE1001118, PLACE1001238, PLACE1001304, PL ACE1001602,
        PLACE1001611, PLACE1001632, PLACE1002450, PLACE1002532, PLACE1002834, P LACE1003100,
        PLACE1003190, PLACE1003302, PLACE1003605, PLACE1003704, PLACE1003738, PLACE1003885,
        PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004681, PLACE1004902, PLACE1005287,
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        PLACE1005876, PLACE1006011, PLACE1006119, PLACE1006414, PLACE100643 8, PLACE1006482,
        PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10079 55, PLACE1007969,
        PLACE1008044, PLACE1008177, PLACE1008603, PLACE1008627,
        PLACE1008941, PLACE1008947, PLACE1009099, PLACE1009113, PLACE1009246, PLACE1009398,
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        PLACE1010152, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010702, PLACE101072 0,
        PLACE1010771, PLACE1010870, PLACE1011056, PLACE1011229, PLACE1011576, PLACE10116 64,
        PLACE2000072, PLACE2000359, PLACE2000411, PLACE4000014, PLACE4000128, PLACE4000 156,
        PLACE4000259, PLACE4000261, PLACE4000431, THYRO1000121, THYRO1000242, THYRO100 0501,
        THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000214, Y79AA10 00539,
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        Y79AA1000589, Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1000784, Y79AA1 001105,
        Y79AA1001299, Y79AA1001391, Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA 1001866,
        Y79AA1001963, Y79AA1002093, Y79AA1002472, Y79AA1002482,
        (11) The following 111 clones were predicted to encode a protein in the category of proteins associated with protein
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        synthesis or transport (including the clones belonging to plural categories).
        HEMBA1000012, HEMBA1000542, HEMBA1001913, HEMBA1003773, HEMBA1004202, HEMBA1004356,
        HEMBA1004847, HEMBA1005047, HEMBA1005202, HEMBA1005963, HEMBA1006310, HEMBA100665 2,
        HEMBA1006914, HEMBB1000217, HEMBB1000725, HEMBB1001346, HEMBB1001831, MAMMA10000 85,
        MAMMA1000734, MAMMA1002170, MAMMA1002236, MAMMA1002598, NT2RM1000661, NT2RM2000 504,
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        NT2RM2000577, NT2RM2001131, NT2RM4000155, NT2RM4001444, NT2RM4002062, NT2RM400 2205,
        NT2RM4002623, NT2RP1000326, NT2RP1000947, NT2RP1001569, NT2RP2000710, NT2RP20 00880,
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NT2RP2001290, NT2RP2001511, NT2RP2001660, NT2RP2002185, NT2RP2002606, NT2RP2 002959, NT2RP2002980, NT2RP2003158, NT2RP2003760, NT2RP2004194, NT2RP2004791, NT2RP 2005012,

NT2RP2005116, NT2RP2005204, NT2RP2006565, NT2RP2006598, NT2RP3000047, NT2R P3000366, NT2RP3000759, NT2RP3000968, NT2RP3001587, NT2RP3002529, NT2RP3003385, NT2 RP3003589, NT2RP3003876, NT2RP3004618, NT2RP4000370, NT2RP4000524, NT2RP4000528, NT 2RP4001041, NT2RP4001574, NT2RP4001592, NT2RP4002047, OVARC1000004, OVARC1000771, O VARC1000862, OVARC1001180, OVARC1001342, OVARC1001766, OVARC1002138, PLACE1000661, PLACE1000492, PLACE1000863, PLACE1001748, PLACE1002090, PLACE1003394, PLACE1003915, PLACE1004149, PLACE1004743, PLACE1005557, PLACE1005813, PLACE1006488, PLACE100653 1, PLACE1006615, PLACE1007706, PLACE1008273, PLACE1008790, PLACE1008813, PLACE10097 21, PLACE1009763, PLACE1009845, PLACE1010074, PLACE1011109, PLACE1011477,

10 PLACE2000404, PLACE300059, PLACE3000121, PLACE4000558, PLACE4000654, SKNMC1000011, THYRO1000288, THYRO1000974, Y79AA1000346, Y79AA1000968, Y79AA1002209,

(12) The following 23 clones were predicted to encode a protein in the category of proteins associated with cellular defense (including the clones belonging to plural categories).

HEMBA1000005, HEMBA1000531, HEMBB1000217, HEMBB1000399, MAMMA1000734, NT2RP1000493, NT2RP1000493, NT2RP2000006, NT2RP2000045, NT2RP2001536, NT2RP2005204, NT2RP300059 0, NT2RP3001426, NT2RP3002056, NT2RP3004262, NT2RP4001638, OVARC1001900, PLACE10056 11, PLACE1006958, PLACE1008275, PLACE1009113, PLACE4000014, Y79AA1002139,

(13) The following 23 clones were predicted to encode a protein in the category of proteins associated with development or growth (including the clones belonging to plural categories).
 HEMBB1001802, HEMBB1002442, NT2RM2002142, NT2RM4001047, NT2RP2003308, NT2RP2004816, NT2RP3000994, NT2RP3001340, NT2RP3004206, NT2RP3004472, NT2RP4000246, NT2RP400100 6, OVARC1000304, OVARC1000746, OVARC1000996, PLACE1006037, PLACE1009622, PLACE10116 64, Y79AA1001692,

Table 9
Selected clones having the maximal ATGpr1 score of 0.3 or higher (3690 clones).

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	clone	name of	maximal
	name	sequence	ATGpr1 score
35		<del></del>	
	HEMBA1000005	F-HEMBA1000005	0.84
	HEMBA1000012	F-HEMBA1000012	0. 56
40	HEMBA1000020	F-HEMBA1000020	0.94
40	HEMBA1000030	F-HEMBA1000030	0. 44
	HEMBA1000046	F-HEMBA1000046	0.50
	HEMBA1000050	F-HEMBA1000050	0.94
45	HEMBA1000076	F-HEMBA1000076	0. 48
	HEMBA1000129	F-HEMBA1000129	0.74
	HEMBA1000141	F-HEMBA1000141	0.55
	HEMBA1000150	F-HEMBA1000150	0.72
50	HEMBA1000156	F-HEMBA1000156	0. 94
	HEMBA1000158	F-HEMBA1000158	0. 62
	HEMBA1000168	F-HEMBA1000168	0.94
	HEMBA1000185	F-HEMBA1000185	0.86
55	HEMBA1000193	F-HEMBA1000193	0.94

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	HEMBA1000201	F-HEMBA1000201	0.83
	HEMBA1000216	F-HEMBA1000216	0, 32
	HEMBA1000227	F-HEMBA1000227	0. 72
5	HEMBA1000231	F-HEMBA1000231	
	HEMBA1000244	F-HEMBA1000244	0.51
	HEMBA1000244	F-HEMBA1000288	0. 36
	HEMBA1000290	F-HEMBA1000290	0. 45
. 10	HEMBA1000302	F-HEMBA1000302	0.71
	HEMBA1000302	F-HEMBA1000303	0. 42
	HEMBA1000304	F-HEMBA1000304	0. 45
	HEMBA1000307	F-HEMBA1000307	0. 48
15	HEMBA1000307	F-HEMBA1000327	0. 94
15	HEMBA1000356	F-HEMBA1000356	0. 90
	HEMBA1000369	F-HEMBA1000369,	0. 94
	HEMBA1000387	F-HEMBA1000387	0. 66
	HEMBA1000392	F-HEMBA1000392	0. 60
20	HEMBA1000396	F-HEMBA1000396	0. 81
	HEMBA1000356	F-HEMBA1000456	0. 90
	HEMBA1000460	F-HEMBA1000460	0.31
	HEMBA1000488	F-HEMBA1000488	0. 47
25	HEMBA1000490	F-HEMBA1000490	0. 89
	HEMBA1000491	F-HEMBA1000491	0. 94
	HEMBA1000501	F-HEMBA1000501	0. 67
	HEMBA1000505	F-HEMBA1000505	0. 65
30	HEMBA1000508	F-HEMBA1000508	0. 39
	HEMBA1000520	F-HEMBA1000520	0. 94
•	HEMBA1000523	F-HEMBA1000523	0. 86
	HEMBA1000531	F-HEMBA1000531	0.49
35	HEMBA1000534	F-HEMBA1000534	0. 52
	HEMBA1000542	F-HEMBA1000542	0.81
	HEMBA1000555	F-HEMBA1000555	0. 67
	HEMBA1000561	F-HEMBA1000561	0.34
40	HEMBA1000568	F-HEMBA1000568	0.66
	HEMBA1000588	F-HEMBA1000588	0.79
	HEMBA1000591	F-HEMBA1000591	0.92
	HEMBA1000592	F-HEMBA1000592	0.73
45	HEMBA1000594	F-HEMBA1000594	0.89
	HEMBA1000608	F-HEMBA1000608	0.94
	HEMBA1000636	F-HEMBA1000636	0.94
	HEMBA1000637	F-HEMBA1000637	0.92
50	HEMBA1000657	F-HEMBA1000657	0.85
50	HEMBA1000682	F-HEMBA1000682	0.94
	HEMBA1000686	F-HEMBA1000686	0.90
	HEMBA1000719	F-HEMBA1000719	0.94
	HEMBA1000727	F-HEMBA1000727	0.40
55	HEMBA1000752	F-HEMBA1000752	0.31

	HEMBA1000817	F-HEMBA1000817	0. 94
	HEMBA1000851	F-HEMBA1000851	0. 63
	HEMBA1000867	F-HEMBA1000867	0. 68
5	HEMBA1000869	F-HEMBA1000869	0.82
	HEMBA1000872	F-HEMBA1000872	0.54
	HEMBA1000910	F-HEMBA1000910	0. 91
	HEMBA1000918	F-HEMBA1000918	0.38
10	HEMBA1000919	F-HEMBA1000919	0.83
	HEMBA1000942	F-HEMBA1000942	0.31
	HEMBA1000946	F-HEMBA1000946	0. 75
	HEMBA1000968	F-HEMBA1000968	0.39
15	HEMBA1000971	F-HEMBA1000971	0.74
	HEMBA1000975	F-HEMBA1000975	0.46
	HEMBA1000986	F-HEMBA1000986,	0.89
	HEMBA1001008	F-HEMBA1001008	0.47
20	HEMBA1001009	F-HEMBA1001009	0.86
20	HEMBA1001022	F-HEMBA1001022	0.57
	HEMBA1001043	F-HEMBA1001043	0.79
	HEMBA1001052	F-HEMBA1001052	0.83
	HEMBA1001059	F-HEMBA1001059	0.78
25	HEMBA1001080	F-HEMBA1001080	0. 77
	HEMBA1001085	F-HEMBA1001085	0.92
	HEMBA1001088	F-HEMBA1001088	0.94
	HEMBA1001109	F-HEMBA1001109	0.50
30	HEMBA1001122	F-HEMBA1001122	0.74
	HEMBA1001133	F-HEMBA1001133	0. 45
	HEMBA1001137	F-HEMBA1001137	0. 37
	HEMBA1001140	F-HEMBA1001140	0.43
35	HEMBA1001174	F-HEMBA1001174	0.44
	HEMBA1001197	F-HEMBA1001197	0.65
	HEMBA1001213	F-HEMBA1001213	0.61
	HEMBA1001235	F-HEMBA1001235	0. 94
40	HEMBA1001247	F-HEMBA1001247	0.31
	HEMBA1001257	F-HEMBA1001257	0. 92
	HEMBA1001281	F-HEMBA1001281	0.94
	HEMBA1001286	F-HEMBA1001286	0.49
45	HEMBA1001289	F-HEMBA1001289	0.94
	HEMBA1001302	F-HEMBA1001302	0.62
	HEMBA1001303	F-HEMBA1001303	0.79
	HEMBA1001310	F-HEMBA1001310	0.66
50	HEMBA1001326	F-HEMBA1001326	0.59
	HEMBA1001351	F-HEMBA1001351	0.65
	HEMBA1001361	F-HEMBA1001361	0.94
	HEMBA1001377	F-HEMBA1001377	0.87
55	HEMBA1001387	F-HEMBA1001387	0.94
	HEMBA1001388	F-HEMBA1001388	0. 57

	HEMBA1001398	F-HEMBA1001398	0.74
	HEMBA1001405	F-HEMBA1001405	0. 61
	HEMBA1001407	F-HEMBA1001407	0. 43
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	HEMBA1001413	F-HEMBA1001413_	0, 40
	HEMBA1001415	F-HEMBA1001415	0.94
	HEMBA1001446	F-HEMBA1001446	0.90
	HEMBA1001450	F-HEMBA1001450	0.40
10	HEMBA1001455	F-HEMBA1001455	0.73
	HEMBA1001476	F-HEMBA1001476	0.65
	HEMBA1001497	F-HEMBA1001497	0.48
	HEMBA1001510	F-HEMBA1001510	0.36
15	HEMBA1001526	F-HEMBA1001526	0.66
	HEMBA1001533	F-HEMBA1001533	0.46
	HEMBA1001569	F-HEMBA1001569,	
	HEMBA1001570	F-HEMBA1001570	0.53
20	HEMBA1001579	F-HEMBA1001579	0.82
	HEMBA1001581	F-HEMBA1001581	0.80
	HEMBA1001595	F-HEMBA1001595	0.72
	HEMBA1001620	F-HEMBA1001620	0.94
25	HEMBA1001635	F-HEMBA1001635	0.87
20	HEMBA1001640	F-HEMBA1001640	0.82
	HEMBA1001647	F-HEMBA1001647	0.91
	HEMBA1001655	F-HEMBA1001655	0.39
	HEMBA1001661	F-HEMBA1001661	0.86
30	HEMBA1001672	F-HEMBA1001672	0.94
	HEMBA1001702	F-HEMBA1001702	0.72
	HEMBA1001711	F-HEMBA1001711	0.74
	HEMBA1001714	F-HEMBA1001714	0.89
35	HEMBA1001723	F-HEMBA1001723	0.94
	HEMBA1001731	F-HEMBA1001731	0.37
	HEMBA1001744	F-HEMBA1001744	0.31
	HEMBA1001746	F-HEMBA1001746	0.39
40	HEMBA1001781	F-HEMBA1001781	0.67
	HEMBA1001800	F-HEMBA1001800	0.69
	HEMBA1001804	F-HEMBA1001804	0.85
	HEMBA1001809	F-HEMBA1001809	0. 78
45	HEMBA1001815	F-HEMBA1001815	0.49
	HEMBA1001819	F-HEMBA1001819	0. 90
	HEMBA1001822	F-HEMBA1001822	0.80
	HEMBA1001824	F-HEMBA1001824	0.68
50	HEMBA1001847	F-HEMBA1001847	0.46
	HEMBA1001864	F-HEMBA1001864	0.34
	HEMBA1001866	F-HEMBA1001866	0.53
	HEMBA1001869	F-HEMBA1001869	0.69
55	HEMBA1001896	F-HEMBA1001896	0. 48
33	HEMBA1001910	F-HEMBA1001910	0.84

	HEMBA1001912	F-HEMBA1001912	0. 48
	HEMBA1001913	F-HEMBA1001913	0. 94
	HEMBA1001915	F-HEMBA1001915	0. 41
5	HEMBA1061921	F-HEMBA1001921	0. 94
	HEMBA1001939	F-HEMBA1001939	0.87
	HEMBA1001950	F-HEMBA1001950	0. 36
	HEMBA1001967	F-HEMBA1001967	0.94
10	HEMBA1001987	F-HEMBA1001987	0.41
	HEMBA1002018	F-HEMBA1002018	0.41
	HEMBA1002035	F-HEMBA1002035	0.36
,	HEMBA1002049	F-HEMBA1002049	0. 39
15	HEMBA1002084	F-HEMBA1002084	0. 45
	HEMBA1002092	F-HEMBA1002092	0.94
	HEMBA1002102	F-HEMBA1002102,	0.94
	HEMBA1002119	F-HEMBA1002119	0.60
20	HEMBA1002125	F-HEMBA1002125	0.88
	HEMBA1002150	F-HEMBA1002150	0. 31
	HEMBA1002151	F-HEMBA1002151	0.82
	HEMBA1002161	F-HEMBA1002161	0.94
25	HEMBA1002177	F-HEMBA1002177	0.38
	HEMBA1002189	F-HEMBA1002189	0.43
	HEMBA1002191	F-HEMBA1002191	0.94
	HEMBA1002199	F-HEMBA1002199	0.85
	HEMBA1002212	F-HEMBA1002212	0.39
30	HEMBA1002215	F-HEMBA1002215	0.94
	HEMBA1002229	F-HEMBA1002229	0.83
	HEMBA1002237	F-HEMBA1002237	0.86
	HEMBA1002241	F-HEMBA1002241	0.66
35	HEMBA1002265	F-HEMBA1002265	0. 38
	HEMBA1002267	F-HEMBA1002267	0. 33
	HEMBA1002341	F-HEMBA1002341	0. 53
	HEMBA1002363	F-HEMBA1002363	0.88
40	HEMBA1002417	F-HEMBA1002417	0. 83
	HEMBA1002419	F-HEMBA1002419	0. 92
	HEMBA1002430	F-HEMBA1002430	0. 45
	HEMBA1002439	F-HEMBA1002439	0.83
45	HEMBA1002458		0.47
	HEMBA1002460		0.48
	HEMBA1002462		0.55
	HEMBA1002469		0.79
50 ·	HEMBA1002475		0.88
	HEMBA1002477		0.45
	HEMBA1002495		0. 78 0. 94
	HEMBA1002503		
55	HEMBA1002508		0.33
	HEMBA1002513	F-HEMBA1002513	0. 41

	HEMBA1002515	F-HEMBA1002515	0.94
	HEMBA1002542	F-HEMBA1002542	0.46
	HEMBA1002547	F-HEMBA1002547	0. 45
5	HEMBA1002569	F-HEMBA1002569	0.86
	HEMBA1002583	F-HEMBA1002583	0.86
	HEMBA1002609	F-HEMBA1002609	0. 75
	HEMBA1002624	F-HEMBA1002624	0. 59
10	HEMBA1002688	F-HEMBA1002688	0.88
	HEMBA1002696	F-HEMBA1002696	0. 82
	HEMBA1002703	F-HEMBA1002703	0.94
	HEMBA1002746	F-HEMBA1002746	0. 33
15	HEMBA1002750	F-HEMBA1002750	0. 53
	HEMBA1002768	F-HEMBA1002768	0. 94
	HEMBA1002770	F-HEMBA1002770,	
	HEMBA1002777	F-HEMBA1002777	0. 90
20	HEMBA1002779	F-HEMBA1002779	0. 92
	HEMBA1002794	F-HEMBA1002794	0.86
	HEMBA1002810	F-HEMBA1002810	0. 87
	HEMBA1002816	F-HEMBA1002816	0.91
25	HEMBA1002818	F-HEMBA1002818	0.94
25	HEMBA1002850	F-HEMBA1002850	0.57
	HEMBA1002863	F-HEMBA1002863	0.48
	HEMBA1002876	F-HEMBA1002876	0.86
22	HEMBA1002935	F-HEMBA1002935	0.91
30	HEMBA1002937	F-HEMBA1002937	0. 61
	HEMBA1002939	F-HEMBA1002939	0.32
	HEMBA1002951	F-HEMBA1002951	0.83
	HEMBA1002954	F-HEMBA1002954	0.67
35	HEMBA1002970	F-HEMBA1002970	0.33
	HEMBA1002971	F-HEMBA1002971	0. 57
	HEMBA1002973	F-HEMBA1002973	0. 33
	HEMBA1002997	F-HEMBA1002997	0. 46
40	HEMBA1002999	F-HEMBA1002999	0.49
	HEMBA1003021	F-HEMBA1003021	0. 74
	HEMBA1003033	F-HEMBA1003033	0. 94
	HEMBA1003035	F-HEMBA1003035	0. 61
45	HEMBA1003041	F-HEMBA1003041	0. 40
	HEMBA1003046	F-HEMBA1003046	0. 94
	HEMBA1003067	F-HEMBA1003067	0. 34
	HEMBA1003077	F-HEMBA1003077	0. 83
50	HEMBA1003078	F-HEMBA1003078	0. 89
·	HEMBA1003079	F-HEMBA1003079	0. 49
	HEMBA1003096	F-HEMBA1003096	0.34
	HEMBA1003117	F-HEMBA1003117	0.94
55	HEMBA1003129	F-HEMBA1003129	0.81
	HEMBA1003136	F-HEMBA1003136	0. 77

	HEMBA1003148	F-HEMBA1003148	0.91
	HEMBA1003175	F-HEMBA1003175	0.41
	HEMBA1003179	F-HEMBA1003179	0.42
5	HEMBA1003199	F-HEMBA1003199	0.84
	HEMBA1003222	F-HEMBA1003222	0.47
	HEMBA1003235	F-HEMBA1003235	0.94
	HEMBA1003250	F-HEMBA1003250	0.39
10	HEMBA1003257	F-HEMBA1003257	0.88
	HEMBA1003281	F-HEMBA1003281	0.67
	HEMBA1003286	F-HEMBA1003286	0.57
	HEMBA1003291	F-HEMBA1003291	0.89
15	HEMBA1003322	F-HEMBA1003322	0.60
	HEMBA1003327	F-HEMBA1003327	0.88
	HEMBA1003369	F-HEMBA1003369,	0.89
	HEMBA1003370	F-HEMBA1003370	0.75
20	HEMBA1003380	F-HEMBA1003380	0.68
	HEMBA1003395	F-HEMBA1003395	0.88
	HEMBA1003402	F-HEMBA1003402	0.45
	HEMBA1003403	F-HEMBA1003403	0.44
25	HEMBA1003408	F-HEMBA1003408	0.34
	HEMBA1003417	F-HEMBA1003417	0.78
	HEMBA1003418	F-HEMBA1003418	0.94
	HEMBA1003433	F-HEMBA1003433	0.70
30	HEMBA1003447	F-HEMBA1003447	0. 49
	HEMBA1003461	F-HEMBA1003461	0. 85
	HEMBA1003463	F-HEMBA1003463	0. 59
	HEMBA1003480	F-HEMBA1003480	0.94
35	HEMBA1003528	F-HEMBA1003528	0. 90
•	HEMBA1003538	F-HEMBA1003538	0. 47
	HEMBA1003545	F-HEMBA1003545	0. 74
	HEMBA1003555	F-HEMBA1003555	0.94
40	HEMBA1003556	F-HEMBA1003556	0.44
40	HEMBA1003560	F-HEMBA1003560	0.40
	HEMBA1003568	F-HEMBA1003568	0. 94
	HEMBA1003569	F-HEMBA1003569	0. 82
	HEMBA1003581	F-HEMBA1003581	0. 83
45	HEMBA1003591	F-HEMBA1003591	0.68
	HEMBA1003615	F-HEMBA1003615 F-HEMBA1003617	0.94
	HEMBA1003617 HEMBA1003621	F-HEMBA1003621	0.88
	HEMBA1003645	F-HEMBA1003645	0. 47 0. 75
50	HEMBA1003646	F-HEMBA1003646	0. 75
	HEMBA1003662	F-HEMBA1003662	0. 61
	HEMBA1003667	F-HEMBA1003667	0. 54
	HEMBA1003679	F-HEMBA1003679	0. 72
55	HEMBA1003680	F-HEMBA1003680	0. 72
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	HEMBA1003684	F-HEMBA1003684	0. 94
	HEMBA1003690	F-HEMBA1003690	0. 93
	HEMBA1003692	F-HEMBA1003692	0.46
5	HEMBA1003711	F-HEMBA1003711	0. 94
	HEMBA1003720	F-HEMBA1003720	0. 94
	HEMBA1003729	F-HEMBA1003729	0.73
	HEMBA1003742	F-HEMBA1003742	0. 94
10	HEMBA1003760	F-HEMBA1003760	0. 75
	HEMBA1003783	F-HEMBA1003783	0. 31
	HEMBA1003799	F-HEMBA1003799	0.70
	HEMBA1003803	F-HEMBA1003803	0. 87
15	HEMBA1003804	F-HEMBA1003804	0. 94
	HEMBA1003805	F-HEMBA1003805	0.94
	HEMBA1003807	F-HEMBA1003807,	0.91
	HEMBA1003827	F-HEMBA1003827	0.62
20	HEMBA1003836	F-HEMBA1003836	0.69
	HEMBA1003838	F-HEMBA1003838	0.80
	HEMBA1003864	F-HEMBA1003864	0.94
	HEMBA1003866	F-HEMBA1003866	0.32
25	HEMBA1003879	F-HEMBA1003879	0.82
	HEMBA1003880	F-HEMBA1003880	0. 59
	HEMBA1003893	F-HEMBA1003893	0. 47
	HEMBA1003953	F-HEMBA1003953	0. 77
30	HEMBA1003959	F-HEMBA1003959	0. 32
30	HEMBA1003978	F-HEMBA1003978	0. 37
	HEMBA1003985	F-HEMBA1003985	0. 31
•	HEMBA1003989	F-HEMBA1003989	0. 75
05	HEMBA1004011	F-HEMBA1004011	0. 34
35	HEMBA1004048	F-HEMBA1004048	0. 32
	HEMBA1004055	F-HEMBA1004055	0.74
	HEMBA1004056	F-HEMBA1004056	0. 49
	HEMBA1004074	F-HEMBA1004074	0. 39
40	HEMBA1004086	F-HEMBA1004086	0. 45
	HEMBA1004097	F-HEMBA1004097	0.82
	HEMBA1004111	F-HEMBA1004111	0.94
	HEMBA1004131	F-HEMBA1004131	0.84
45	HEMBA1004133	F-HEMBA1004133	0.40
	HEMBA1004143	F-HEMBA1004143	0.33
	HEMBA1004146	F-HEMBA1004146	0.31
	HEMBA1004150	F-HEMBA1004150	0.70
50	HEMBA1004168	F-HEMBA1004168	0.34
	HEMBA1004199	F-HEMBA1004199	0.34
	HEMBA1004200	F-HEMBA1004200 F-HEMBA1004202	0. 40 0. 88
	HEMBA1004202 HEMBA1004203	F-HEMBA1004202	0.88
55	HEMBA1004203	F-HEMBA1004207	0. 53
	FILMOA 1 UU420 /	FTREMORIUU4ZU/	0.53

	HEMBA1004227	F-HEMBA1004227	0.94
	HEMBA1004238	F-HEMBA1004238	0.44
	HEMBA1004246	F-HEMBA1004246	0.71
5	HEMBA1004248	F-HEMBA1004248	0.40
	HEMBA1004264	F-HEMBA1004264	0. 94
	HEMBA1004274	F-HEMBA1004274	0.88
	HEMBA1004275	F-HEMBA1004275	0. 85
10	HEMBA1004276	F-HEMBA1004276	0. 32
	HEMBA1004286	F-HEMBA1004286	0.37
	HEMBA1004289	F-HEMBA1004289	0.76
	HEMBA1004321	F-HEMBA1004321	0.82
15	HEMBA1004327	F-HEMBA1004327	0.40
	HEMBA1004330	F-HEMBA1004330	0.84
	HEMBA1004335	F-HEMBA1004335,	0. 79
	HEMBA1004341	F-HEMBA1004341	0. 38
20	HEMBA1004353	F-HEMBA1004353	0.79
	HEMBA1004372	F-HEMBA1004372	0. 94
	HEMBA1004389	F-HEMBA1004389	0. 32
	HEMBA1004408	F-HEMBA1004408	0.70
25	HEMBA1004479	F-HEMBA1004479	0.55
	HEMBA1004499	F-HEMBA1004499	0. 62
•	HEMBA1004502	F-HEMBA1004502	0. 31
	HEMBA1004507	F-HEMBA1004507	0.71
30	HEMBA1004509	F-HEMBA1004509	0. 68
	HEMBA1004534	F-HEMBA1004534	0.56
	HEMBA1004542	F-HEMBA1004542	0.36
	HEMBA1004554	F-HEMBA1004554	0. 90
35	HEMBA1004560	F-HEMBA1004560	0. 57
	HEMBA1004573	F-HEMBA1004573	0. 93
	HEMBA1004596	F-HEMBA1004596	0. 45
	HEMBA1004604	F-HEMBA1004604	0. 91
40	HEMBA1004610	F-HEMBA1004610	0. 59
•	HEMBA1004632	F-HEMBA1004632	0. 59
	HEMBA1004637	F-HEMBA1004637	0. 38
	HEMBA1004638	F-HEMBA1004638	0.40
45	HEMBA1004669	F-HEMBA1004669	0. 94
45	HEMBA1004693	F-HEMBA1004693	0. 52
	HEMBA1004697	F-HEMBA1004697	0.94
	HEMBA1004705 HEMBA1004709	F-HEMBA1004705 F-HEMBA1004709	0.94
	HEMBA1004709	F-HEMBA1004711	0. 77 0. 73
50		F-HEMBA1004711	0.73
	HEMBA1004725 HEMBA1004734	F-HEMBA1004734	0. 94
	HEMBA1004734	F-HEMBA1004736	0. 94
	HEMBA1004751	F-HEMBA1004751	0. 64
55	HEMBA1004751	F-HEMBA1004752	0. 87
	11LINDA ( 004 / 32		0.07

	HEMBA1004753	F-HEMBA1004753 ·	0. 54
	HEMBA1004756	F-HEMBA1004756	0. 54
	HEMBA1004758	F-HEMBA1004758	0. 69
5	HEMBA1004763	F-HEMBA1004763	
	HEMBA1004768	F-HEMBA1004768	0. 66
	HEMBA1004771	F-HEMBA1004771	0. 72
	HEMBA1004776	F-HEMBA1004776	0.49
10	HEMBA1004795	F-HEMBA1004795	0.46
	HEMBA1004806	F-HEMBA1004806	0.32
	HEMBA1004847	F-HEMBA1004847	0.79
	HEMBA1004850	F-HEMBA1004850	0.94
15	HEMBA1004863	F-HEMBA1004863	0. 32
	HEMBA1004864	F-HEMBA1004864	0.94
	HEMBA1004889	F-HEMBA1004889,	0.94
	HEMBA1004923	F-HEMBA1004923	0.70
20	HEMBA1004929		0.31
•	HEMBA1004930	F-HEMBA1004930	0.90
	HEMBA1004933	F-HEMBA1004933	0.84
	HEMBA1004934	F-HEMBA1004934	0.34
25	HEMBA1004944	F-HEMBA1004944	0.40
25	HEMBA1004954	F-HEMBA1004954	0.62
	HEMBA1004972	F-HEMBA1004972	0.90
	HEMBA1004973	F-HEMBA1004973	0.94
00	HEMBA1004977	F-HEMBA1004977	0.77
30	HEMBA1004980	F-HEMBA1004980	0.33
	HEMBA1005009	F-HEMBA1005009	0.94
	HEMBA1005019	F-HEMBA1005019	0.32
	HEMBA1005029	F-HEMBA1005029	0.84
35	HEMBA1005035	F-HEMBA1005035	0.91
	HEMBA1005047	F-HEMBA1005047	0.94
	HEMBA1005050	F-HEMBA1005050	0.88
	HEMBA1005066	F-HEMBA1005066	0.79
40	HEMBA1005075	F-HEMBA1005075	0.31
	HEMBA1005079	F-HEMBA1005079	0.35
	HEMBA1005083	F-HEMBA1005083	0. 43
	HEMBA1005101	F-HEMBA1005101	0.94
45	HEMBA1005113	F-HEMBA1005113	0.38
	HEMBA1005133	F-HEMBA1005133	0.34
	HEMBA1005149	F-HEMBA1005149	0. 36
	HEMBA1005185	F-HEMBA1005185	0.64
50	HEMBA1005201	F-HEMBA1005201	0.83
	HEMBA1005202	F-HEMBA1005202	0. 94
	HEMBA1005206	F-HEMBA1005206	0.94
	HEMBA1005219	F-HEMBA1005219	0. 57
55	HEMBA1005223	F-HEMBA1005223	0. 33
	HEMBA1005244	F-HEMBA1005244	0. 35

	HEMBA1005252	F-HEMBA1005252	0. 80
	HEMBA1005296	F-HEMBA1005296	0. 94
	HEMBA1005314	F-HEMBA1005314	0. 94
<b>5</b> .	HEMBA1005331	F-HEMBA1005331	0. 69
	HEMBA1005338	F-HEMBA1005338	0. 94
	HEMBA1005367	F-HEMBA1005367	0. 50
	HEMBA1005382	F-HEMBA1005382	0. 91
10	HEMBA1005394	F-HEMBA1005394	0, 45
	HEMBA1005403	F-HEMBA1005403	0.94
•	HEMBA1005411	F-HEMBA1005411	0. 33
	HEMBA1005423	F-HEMBA1005423	0. 64
15	HEMBA1005468	F-HEMBA1005468	0. 33
-	HEMBA1005469	F-HEMBA1005469	0. 59
	HEMBA1005472	F-HEMBA1005472,	
	HEMBA1005474	F-HEMBA1005474	0. 51
20	HEMBA1005475	F-HEMBA1005475	0. 36
20	HEMBA1005513	F-HEMBA1005513	0.94
	HEMBA1005517	F-HEMBA1005517	0. 94
	HEMBA1005518	F-HEMBA1005518	0.39
	HEMBA1005526	F-HEMBA1005526	0. 62
25	HEMBA1005528	F-HEMBA1005528	0.81
	HEMBA1005530	F-HEMBA1005530	0. 62
	HEMBA1005548	F-HEMBA1005548	0.94
	HEMBA1005558	F-HEMBA1005558	0.94
30	HEMBA1005576	F-HEMBA1005576	0.49
	HEMBA1005581	F-HEMBA1005581	0.53
	HEMBA1005582	F-HEMBA1005582	0.34
	HEMBA1005583	F-HEMBA1005583	0. 94
35	HEMBA1005595	F-HEMBA1005595	0. 93
	HEMBA1005609	F-HEMBA1005609	0. 34
	HEMBA1005621	F-HEMBA1005621	0.94
	HEMBA1005666	F-HEMBA1005666	0. 57
40	HEMBA1005680	F-HEMBA1005680	0. 38
	HEMBA1005685	F-HEMBA1005685	0. 32
	HEMBA1005732	F-HEMBA1005732	0. 33
	HEMBA1005737	F-HEMBA1005737	0. 92
45	HEMBA1005746	F-HEMBA1005746	0. 37
	HEMBA1005755	F-HEMBA1005755	0.94
	HEMBA1005813	F-HEMBA1005813	0. 33
	HEMBA1005815	F-HEMBA1005815	0. 82
50	HEMBA1005822	F-HEMBA1005822	0. 83
	HEMBA1005834	F-HEMBA1005834	0. 70
	HEMBA1005852	F-HEMBA1005852	0. 76
	HEMBA1005884	F-HEMBA1005884	0. 79
55	HEMBA1005891	F-HEMBA1005891	0. 94
	HEMBA1005963	F-HEMBA1005963	0. 40

	HEMBA1005990 F-HEMBA1005990 (	). 82
	HEMBA1005991 F-HEMBA1005991 (	0. 43
	HEMBA1006031 F-HEMBA1006031 (	0. 55
5	HEMBA1006035 F-HEMBA1006035 (	0. 43
	HEMBA1006081 F-HEMBA1006081 (	0. 36
	HEMBA1006091 F-HEMBA1006091 (	0. 81
	HEMBA1006100 F-HEMBA1006100 (	0. 31
10	HEMBA1006108 F-HEMBA1006108 (	0. 80
		0. 45
	HEMBA1006130 F-HEMBA1006130 (	0. 46
	HEMBA1006138 F-HEMBA1006138 (	0. 48
15		0. 72
	HEMBA1006173 F-HEMBA1006173	0. 42
	HEMBA1006182 F-HEMBA1006182,	0. 73
	·	0. 94
20	HEMBA1006235 F-HEMBA1006235	0. 58
20	HEMBA1006248 F-HEMBA1006248	0. 57
	HEMBA1006252 F-HEMBA1006252	0. 31
	HEMBA1006272 F-HEMBA1006272	0. 54
0.5	HEMBA1006278 F-HEMBA1006278	0. 92
25	HEMBA1006283 F-HEMBA1006283	0. 90
	HEMBA1006284 F-HEMBA1006284	0.93
	HEMBA1006291 F-HEMBA1006291	0. 38
	HEMBA1006293 F-HEMBA1006293	0.89
30		0.94
	***************************************	0. 67
		0. 94
		0. 94
35	***************************************	0. 56
		0. 94
		0. 33
		0.71
40		0.40
	HEMBA1006445 F-HEMBA1006445	0. 86
	HEMBA1006467 F-HEMBA1006467	0. 55
	HEMBA1006474 F-HEMBA1006474	0. 66
45	HEMBA1006483 F-HEMBA1006483	0. 71
	HEMBA1006485 F-HEMBA1006485	0. 87
	HEMBA1006492 F-HEMBA1006492	0. 73
	HEMBA1006494 F-HEMBA1006494	0. 90
50	HEMBA1006497 F-HEMBA1006497	0. 77
	HEMBA1006502 F-HEMBA1006502	0. 80
	HEMBA1006507 F-HEMBA1006507	0. 56
	HEMBA1006521 F-HEMBA1006521	0.91
55	HEMBA1006530 F-HEMBA1006530	0. 37
	HEMBA1006535 F-HEMBA1006535	056

	HEMBA1006559	F-HEMBA1006559	0. 39
	HEMBA1006566	F-HEMBA1006566	0. 41
	HEMBA1006579	F-HEMBA1006579	0. 83
5	• • • • • • • • • • • • • • • • • • • •		
5	HEMBA1006583	F-HEMBA1006583	0.31
	HEMBA1006612	F-HEMBA1006612	0. 55
	HEMBA1006624	F-HEMBA1006624	0. 94
	HEMBA1006631	F-HEMBA1006631	0. 74
10	HEMBA1006643	F-HEMBA1006643	0.38
	HEMBA1006652	F-HEMBA1006652	0. 93
	HEMBA1006659	F-HEMBA1006659	0. 63
	HEMBA1006674	F-HEMBA1006674	0.54
15	HEMBA1006682	F-HEMBA1006682	0.34
	HEMBA1006708	F-HEMBA1006708	0.40
	HEMBA1006709	F-HEMBA1006709,	0.41
	HEMBA1006717	F-HEMBA1006717	0. 43
20	HEMBA1006737	F-HEMBA1006737	0. 72
	HEMBA1006754	F-HEMBA1006754	·0. 45
	HEMBA1006758	F-HEMBA1006758	0.60
	HEMBA1006767	F-HEMBA1006767	0.61
25	HEMBA1006789	F-HEMBA1006789	0.34
	HEMBA1006795	F-HEMBA1006795	0.48
	HEMBA1006796	F-HEMBA1006796	0. 41
	HEMBA1006807	F-HEMBA1006807	0.34
	HEMBA1006832	F-HEMBA1006832	0. 59
30	HEMBA1006877	F-HEMBA1006877	0.90
	HEMBA1006885	F-HEMBA1006885	0.80
	HEMBA1006900	F-HEMBA1006900	0. 34
	HEMBA1006914	F-HEMBA1006914	0. 78
35	HEMBA1006926	F-HEMBA1006926	0.67
	HEMBA1006936	F-HEMBA1006936	0. 65
	HEMBA1006941	F-HEMBA1006941	0.94
	HEMBA1006973	F-HEMBA1006973	0.64
40	HEMBA1006976	F-HEMBA1006976	0.40
	HEMBA1006993	F-HEMBA1006993	0.81
	HEMBA1006996	F-HEMBA1006996	0.77
	HEMBA1007002	F-HEMBA1007002	0.72
45	HEMBA1007018	F-HEMBA1007018	0.31
,	HEMBA1007052	F-HEMBA1007052	0.41
	HEMBA1007062	F-HEMBA1007062	0. 39
•	HEMBA1007066	F-HEMBA1007066	0. 35
50	HEMBA1007080	F-HEMBA1007080	0. 43
	HEMBA1007085	F-HEMBA1007085	0. 73
	HEMBA1007087	F-HEMBA1007087	0.94
	HEMBA1007112	F-HEMBA1007112	0. 49
55	HEMBA1007121	F-HEMBA1007121	0. 70
	HEMBA1007149	F-HEMBA1007149	0. 94

	HEMBA1007151	F-HEMBA1007151	0. 80
	HEMBA1007174	F-HEMBA1007174	0. 63
	HEMBA1007178	F-HEMBA1007178	0. 47
5	HEMBA1007194	F-HEMBA1007194	0. 94
	HEMBA1007203	F-HEMBA1007203	0. 48
	HEMBA1007224	F-HEMBA1007224	0. 74
	HEMBA1007243	F-HEMBA1007243	0. 94
10	HEMBA1007251	F-HEMBA1007251	0. 79
	HEMBA1007267	F-HEMBA1007267	0. 86
	HEMBA1007281	F-HEMBA1007281	0. 61
	HEMBA1007300	F-HEMBA1007300	0. 87
15	HEMBA1007301	F-HEMBA1007301	0. 38
73	HEMBA1007301	F-HEMBA1007319	0. 47
	HEMBA1007319	F-HEMBA1007320,	0. 69
	HEMBA1007342	F-HEMBA1007342	0.80
	HEMBB1000008	F-HEMBB1000008	0. 39
20	HEMBB1000018	F-HEMBB1000018	0. 79
	HEMBB100004	F-HEMBB1000024	0. 62
	HEMBB1000025	F-HEMBB1000025	0.66
	HEMBB1000030	F-HEMBB1000030	0. 33
25	HEMBB1000036	F-HEMBB1000036	0.75
	HEMBB1000037	F-HEMBB1000037	0. 94
	HEMBB1000048	F-HEMBB1000048	0.87
	HEMBB1000083	F-HEMBB1000083	0.48
30	HEMBB1000103	F-HEMBB1000103	0. 63
	HEMBB1000119	F-HEMBB1000119	0. 94
•	HEMBB1000136	F-HEMBB1000136	0. 92
	HEMBB1000173	F-HEMBB1000173	0. 46
35	HEMB81000175	F-HEMBB1000175	0.48
	HEMBB1000198	F-HEMBB1000198	0.74
	HEMBB1000215	F-HEMBB1000215	0.61
	HEMBB1000217	F-HEMBB1000217	0. 94
40	HEMBB1000226	F-HEMBB1000226	0.94
·	HEMBB1000240	F-HEMBB1000240	0.40
	HEMBB1000244	F-HEMBB1000244	0.81
	HEMBB1000266	F-HEMBB1000266	0. 92
45	HEMBB1000338	F-HEMB81000338	0. 33
	HEMBB1000339	F-HEMBB1000339	0.40
	HEMBB1000391	F-HEMBB1000391	0.43
	HEMBB1000438	F-HEMB81000438	0.79
50	HEMBB1000449	F-HEMBB1000449	0.48
	HEMBB1000510	F-HEMBB1000510	0. 35
	HEMBB1000550	F-HEMBB1000550	0. 94
	HEMBB1000556	F-HEMBB1000556	0.89
55	HEMBB1000589	F-HEMBB1000589	0. 77
55	HEMBB1000591	F-HEMBB1000591	0. 33

	HEMBB1000592	F-HEMBB1000592	0.77
	HEMBB1000593	F-HEMBB1000593	0.68
	HEMBB1000623	F-HEMBB1000623	0.94
5	HEMBB1000630	F-HEMBB1000630	0.59
	HEMBB1000631	F-HEMBB1000631	0.71
	HEMBB1000632	F-HEMBB1000632	0.38
	HEMBB1000671	F-HEMBB1000671	0.36
10	HEMBB1000673	F-HEMBB1000673	0.54
	HEMBB1000693	F-HEMBB1000693	0.68
	HEMBB1000705	F-HEMBB1000705	0.44
	HEMBB1000706	F-HEMBB1000706	0.45
15	HEMBB1000725	F-HEMBB1000725	0.91
	HEMBB1000763	F-HEMBB1000763	0.87
	HEMBB1000774	F-HEMBB1000774	0.80
	HEMBB1000781	F-HEMBB1000781	0.94
	HEMBB1000789	F-HEMBB1000789	0.34
20	HEMBB1000807	F-HEMBB1000807	0.94
	HEMBB1000810	F-HEMBB1000810	0.33
	HEMBB1000826	F-HEMBB1000826	0.36
	HEMBB1000835	F-HEMBB1000835	0.81
25	HEMBB1000848	F-HEMBB1000848	0.66
	HEMBB1000852	F-HEMB81000852	0.85
	HEMBB1000870	F-HEMB81000870	0.34
	HEMBB1000887	F-HEMBB1000887	0.58
30	HEMB81000908	F-HEMBB1000908	0.31
	HEMBB1000927	F-HEMBB1000927	0.94
	HEMBB1000947	F-HEMBB1000947	0.33
	HEMBB1000973	F-HEMBB1000973	0.72
35	HEMBB1000975	F-HEMBB1000975	0.68
	HEMB81000985	F-HEMBB1000985	0.62
	HEMB81000991	F-HEMB81000991	0. 45
	HEMBB1001011	F-HEMBB1001011	0.58
40	HEMBB1001014	F-HEMBB1001014	0.86
	HEMBB1001020	F-HEMBB1001020	0.87
	HEMB81001024	F-HEMBB1001024	0.64
	HEMBB1001056	F-HEMBB1001056	0.94
45	HEMBB1001058	F-HEMBB1001058	0.39
	HEMB81001068	F-HEMBB1001068	0. 57
	HEMBB1001096	F-HEMBB1001096	0. 48
•	HEMBB1001105	F-HEMBB1001105	0. 45
50	HEMBB1001112	F-HEMBB1001112	0. 47
	HEMBB1001117	F-HEMBB1001117	0. 83
	HEMBB1001126	F-HEMBB1001126	0.76
	HEMBB1001137	F-HEMBB1001137	0. 53
55	HEMBB1001151	F-HEMBB1001151	0. 94
55	HEMBB1001153	F-HEMBB1001153	0.77

	HEMBB1001169	F-HEMBB1001169	0. 39
	HEMBB1001175	F-HEMBB1001175	0. 91
	HEMBB1001182	F-HEMBB1001182	0. 94
5	HEMBB1001199	F-HEMBB1001199	0, 54
	HEMBB1001210	F-HEMBB1001210	0. 88
	HEMBB1001234	F-HEMBB1001234	0. 67
	HEMBB1001242	F-HEMBB1001242	0. 33
10	HEMBB1001288	F-HEMBB1001288	0. 33
	HEMBB1001289	F-HEMBB1001289	0. 58
	HEMBB1001294	F-HEMBB1001294	0. 86
	HEMBB1001314	F-HEMBB1001314	0. 76
15	HEMBB1001331	F-HEMBB1001331	0. 44
	HEMBB1001339	F-HEMBB1001339	0. 40
	HEMBB1001346	F-HEMBB1001346	0. 83
	HEMBB1001364	F-HEMBB1001364	0. 69
20	HEMBB1001369	F-HEMBB1001369	0. 47
20		- F-HEMBB1001384	0. 80
	HEMBB1001387	F-HEMBB1001387	0. 36
	HEMBB1001394	F-HEMBB1001394	0.94
	HEMBB1001410	F-HEMBB1001410	0. 51
25	HEMBB1001426	F-HEMBB1001426	0. 80
	HEMBB1001429	F-HEMBB1001429	0. 91
	HEMBB1001449	F-HEMBB1001449	0. 31
	HEMBB1001482	F-HEMBB1001482	0. 44
30	HEMB81001531	F-HEMBB1001531	0. 58
	HEMBB1001562	F-HEMBB1001562	0.65
	HEMBB1001564	F-HEMBB1001564	0.46
	HEMBB1001585	F-HEMBB1001585	0. 74
35	HEMBB1001603	F-HEMBB1001603	0.81
	HEMBB1001635	F-HEMBB1001635	0. 46
	HEMBB1001653	F-HEMBB1001653	0. 33
	HEMBB1001665	F-HEMBB1001665	0. 39
40	HEMBB1001668	F-HEMBB1001668	0. 38
	HEMBB1001673	F-HEMBB1001673	0. 83
	HEMBB1001684	F-HEMBB1001684	0. 44
	HEMBB1001685	F-HEMBB1001685	0. 90
45	HEMBB1001695	F-HEMBB1001695	0. 92
	HEMBB1001707	F-HEMBB1001707	0. 51
	HEMBB1001736	F-HEMBB1001736	0. 94
	HEMBB1001760	F-HEMBB1001760	0. 88
50	HEMBB1001785	F-HEMBB1001785	0. 74
	HEMBB1001812	F-HEMBB1001812	0. 33
	HEMBB1001816	F-HEMBB1001816	0.41
	HEMBB1001831	F-HEMBB1001831	0. 93
55	HEMBB1001834	F-HEMBB1001834	0. 78
	HEMBB1001839	F-HEMBB1001839	0. 32

	HEMBB1001869	F-HEMBB1001869	0. 48
	HEMBB1001872	F-HEMBB1001872	0.93
	HEMBB1001874	F-HEMBB1001874	0.31
5	HEMBB1001905	F-HEMBB1001905	0. 87
	HEMBB1001906	F-HEMBB1001906	0.88
	HEMBB1001908	F-HEMBB1001908	0.69
	HEMBB1001910	F-HEMBB1001910	0.77
10	HEMBB1001915	F-HEMBB1001915	0.76
	HEMBB1001945	F-HEMBB1001945	0.64
	HEMBB1001947	F-HEMBB1001947	0.94
	HEMBB1001950	F-HEMBB1001950	0.73
15	HEMBB1001957	F-HEMBB1001957	0.64
	HEMBB1001962	F-HEMBB1001962	0.90
	HEMBB1001990	F-HEMBB1001990,	0.83
, i	HEMBB1002044	F-HEMBB1002044	0.82
20	HEMBB1002050	F-HEMBB1002050	0.91
	HEMBB1002068	F-HEMBB1002068	0.48
	HEMBB1002134	F-HEMBB1002134	0.80
	HEMBB1002142	F-HEMBB1002142	0.39
25	HEMBB1002152	F-HEMBB1002152	0.31
23	HEMBB1002193	F-HEMBB1002193	0.35
	HEMBB1002217	F-HEMBB1002217	0.49
	HEMBB1002218	F-HEMBB1002218	0.32
20	HEMBB1002249	F-HEMBB1002249	0.72
30	HEMBB1002300	F-HEMBB1002300	0. 75
	HEMBB1002327	F-HEMBB1002327	0.86
	HEMBB1002329	F-HEMBB1002329	0.31
	HEMBB1002340	F-HEMBB1002340	0. 42
35	HEMBB1002342	F-HEMBB1002342	0. 76
	HEMBB1002358	F-HEMBB1002358	0. 70
	HEMBB1002359	F-HEMBB1002359	0.75
	HEMBB1002383	F-HEMBB1002383	0. 78
40	HEMBB1002409	F-HEMBB1002409	0. 52
	HEMBB1002415	F-HEMBB1002415	0.82
	HEMBB1002442	F-HEMBB1002442	0. 31
	HEMBB1002457	F-HEMBB1002457	0.46
45	HEMBB1002489	F-HEMBB1002489	0.69
	HEMBB1002492	F-HEMB81002492	0.68
	HEMBB1002495	F-HEMBB1002495	0.94
	HEMBB1002502	F-HEMB81002502	0. 39 0. 94
50	HEMBB1002510	F-HEMB81002510	
	HEMBB1002545 HEMBB1002550	F-HEMBB1002545 F-HEMBB1002550	0. 44 0. 57
	HEMBB1002600	F-HEMBB1002600	0. 91
	HEMBB1002607	F-HEMBB1002607	0.83
55	HEMBB1002614	F-HEMBB1002614	0. 75
	110001002014	11LHDD   UUZU   4	0.73

	HEMBB1002684	F-HEMBB1002684	0. 84
	HEMBB1002692	F-HEMBB1002692	0. 52
	HEMBB1002697	F-HEMBB1002697	0. 86
5	HEMBB1002705	F-HEMBB1002705	0. 32
	MAMMA1000019	F-MAMMA1000019	0. 48
	MAMMA1000020	F-MAMMA1000020	0. 36
	MAMMA1000025	F-MAMMA1000025	0. 31
10	MAMMA1000055	F-MAMMA1000055	0. 39
	MAMMA1000057	F-MAMMA1000057	0. 35
	MAMMA1000069	F-MAMMA1000069	0. 78
	MAMMA1000084	F-MAMMA1000084	0. 61
15	MAMMA1000085	F-MAMMA1000085	0.72
	MAMMA1000139	F-MAMMA1000139	0. 33
	MAMMA1000143	F-MAMMA1000143,	0. 40
	MAMMA1000163	F-MAMMA1000163	0. 94
20	MAMMA1000171	F-MAMMA1000171	0. 78
20	MAMMA1000173	F-MAMMA1000173	0. 94
	MAMMA1000183	F-MAMMA1000183	0.77
	MAMMA1000251	F-MAMMA1000251	0. 94
	MAMMA1000257	F-MAMMA1000257	0. 75
25	MAMMA1000277	F-MAMMA1000277	0.94
	MAMMA1000278	F-MAMMA1000278	0. 35
	MAMMA1000279	F-MAMMA1000279	0.62
	MAMMA1000284	F-MAMMA1000284	0. 35
30	MAMMA1000309	F-MAMMA1000309	0.53
	MAMMA1000312	F-MAMMA1000312	0.68
	MAMMA1000313	F-MAMMA1000313	0. 32
	MAMMA1000339	F-MAMMA1000339	0. 35
35	MAMMA1000361	F-MAMMA1000361	0. 61
	MAMMA1000372	F-MAMMA1000372	0.60
	MAMMA1000388	F-MAMMA1000388	0. 93
	MAMMA1000395	F-MAMMA1000395	0.94
40	MAMMA1000410	F-MAMMA1000410	0. 94
	MAMMA1000416	F-MAMMA1000416	0.89
	MAMMA1000421	F-MAMMA1000421	0.50
	MAMMA1000422	F-MAMMA1000422	0.92
45	MAMMA1000429	F-MAMMA1000429	0. 92
	MAMMA1000458	F-MAMMA1000458	0.88
	MAMMA1000468	F-MAMMA1000468	0.60
	MAMMA1000472	F-MAMMA1000472	0. 33
50	MAMMA1000490	F-MAMMA1000490	0.34
	MAMMA1000524	F-MAMMA1000524	0. 39
	MAMMA1000567	F-MAMMA1000567	0.40
	MAMMA1000583	F-MAMMA1000583	0.45
55	MAMMA1000612	F-MAMMA1000612	0.89
	MAMMA1000623	F-MAMMA1000623	0. 44

	MAMMA1000625	F-MAMMA1000625	0. 52
	MAMMA1000664	F-MAMMA1000664	0. 42
	MAMMA1000670	F-MAMMA1000670	0.94
5	MAMMA1000672	F-MAMMA1000672	0. 44
	MAMMA1000684	F-MAMMA1000684	0. 94
	MAMMA1000713	F-MAMMA1000713	0. 88
	MAMMA1000731	F-MAMMA1000731	0. 67
10	MAMMA1000734	F-MAMMA1000734	0.40
	MAMMA1000738	F-MAMMA1000738	0. 93
	MAMMA1000746	F-MAMMA1000746	0.48
	MAMMA1000760	F-MAMMA1000760	0. 94
15	MAMMA1000776	F-MAMMA1000776	0. 94
	MAMMA1000824	F-MAMMA1000824	0. 81
	MAMMA1000831	F-MAMMA1000831,	0. 39
	MAMMA1000841	F-MAMMA1000841	0.41
20	MAMMA1000842	F-MAMMA1000842	0.61
	MAMMA1000843	F-MAMMA1000843	0.48
	MAMMA1000856	F-MAMMA1000856	0.31
	MAMMA1000859	F-MAMMA1000859	0.44
25	MAMMA1000865	F-MAMMA1000865	0. 94
	MAMMA1000875	F-MAMMA1000875	0. 49
	MAMMA1000897	F-MAMMA1000897	0.94
	MAMMA1C00906	F-MAMMA1000906	0. 45
30	MAMMA1000908	F-MAMMA1000908	0.39
	MAMMA1000914	F-MAMMA1000914	0.94
	MAMMA1000921	F-MAMMA1000921	0. 34
	MAMMA1000956	F-MAMMA1000956	0.39
35	MAMMA1000968	F-MAMMA1000968	0. 35
33	MAMMA1000979	F-MAMMA1000979	0.68
	MAMMA1001008	F-MAMMA1001008	0.36
	MAMMA1001021	F-MAMMA1001021	0. 48 0. 34
40	MAMMA1001041 MAMMA1001059	F-MAMMA1001041 F-MAMMA1001059	0. 94
40	MAMMA1001059	F-MAMMA1001059	0. 94
	MAMMA1001075	F-MAMMA1001075	0. 94
	MAMMA1001078	F-MAMMA1001078	0. 52
	MAMMA1001080	F-MAMMA1001080	0. 87
45	MAMMA1001091	F-MAMMA1001091	0.53
	MAMMA1001105	F-MAMMA1001105	0. 94
	MAMMA1G01110	F-MAMMA1001110	0. 78
	MAMMA1001126	F-MAMMA1001126	0.85
50	MAMMA1001139	F-MAMMA1001139	0.38
	MAMMA1001143	F-MAMMA1001143	0. 79
	MAMMA1001154	F-MAMMA1001154	0.31
	MAMMA1001181	F-MAMMA1001181	0. 69
55	MAMMA1001198	F-MAMMA1001198	0.88
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	MAMMA1001202	F-MAMMA1001202	0. 38
	MAMMA1001215	F-MAMMA1001215	0. 94
	MAMMA1001222	F-MAMMA1001213	0. 89
5	MAMMA1001244	F-MAMMA1001244	0. 89
	MAMMA1001259	F-MAMMA1001244	
			0.91
	MAMMA1001260	F-MAMMA1001260	0.75
10	MAMMA1001271	F-MAMMA1001271	0.77
	MAMMA1001292	F-MAMMA1001292	0.94
	MAMMA1001296	F-MAMMA1001296	0.75
	MAMMA1001305	F-MAMMA1001305	0.66
15	MAMMA1001343	F-MAMMA1001343	0.38
15	MAMMA1001346	F-MAMMA1001346	0.81
	MAMMA1001388	F-MAMMA1001388	0. 94
	MAMMA1001411	F-MAMMA1001411,	
	MAMMA1001419	F-MAMMA1001419	0. 45
20	MAMMA1001465	F-MAMMA1001465	0.94
	MAMMA1001476	F-MAMMA1001476	0. 90
	MAMMA1001487	F-MAMMA1001487	0. 78
	MAMMA1001510	F-MAMMA1001510	0. 82
25	MAMMA1001522	F-MAMMA1001522	0.63
	MAMMA1001551	F-MAMMA1001551	0. 81
	MAMMA1001576	F-MAMMA1001576	0.94
	MAMMA1001600	F-MAMMA1001600	0.72
30	MAMMA1001604	F-MAMMA1001604	0.87
	MAMMA1001620	F-MAMMA1001620	0. 43
	MAMMA1001627	F-MAMMA1001627	0. 39
	MAMMA1001630	F-MAMMA1001630	0. 46
35	MAMMA1001633	F-MAMMA1001633	0. 58
33	MAMMA1001635	F-MAMMA1001635	0.89
	MAMMA1001649	F-MAMMA1001649	0. 62
	MAMMA1001654	F-MAMMA1001654	0.94
	MAMMA1001692	F-MAMMA1001692	0.83
40	MAMMA1001730	F-MAMMA1001730	0. 48
	MAMMA1001735	F-MAMMA1001735	0.94
·	MAMMA1001743	F-MAMMA1001743	0. 94
		F-MAMMA1001744	0. 33
45	MAMMA1001751	F-MAMMA1001751	0.94
	MAMMA1001754	F-MAMMA1001754	0. 47
	MAMMA1001757	F-MAMMA1001757	0. 38
	MAMMA1001764	F-MAMMA1001764	0.57
50	MAMMA1001768	F-MAMMA1001768	0.85
	MAMMA1001771	F-MAMMA1001771	0. 39
	MAMMA1001783	F-MAMMA1001783	0.61
	MAMMA1001785	F-MAMMA1001785	0.41
55	MAMMA1001790	F-MAMMA1001790	0. 69
	MAMMA1001812	F-MAMMA1001812	0.32

	MAMMA1001817	F-MAMMA1001817	0, 55
	MAMMA1001824	F-MAMMA1001824	0.57
	MAMMA1001837	F-MAMMA1001837	0.35
5	MAMMA1001848	F-MAMMA1001848	
	MAMMA1001851	F-MAMMA1001851	0.46
	MAMMA1001858	F-MAMMA1001858	0.53
	MAMMA1001864	F-MAMMA1001864	0.32
10	MAMMA1001868	F-MAMMA1001868	0.48
	MAMMA1001874	F-MAMMA1001874	0, 94
	MAMMA1001956	F-MAMMA1001956	0.47
	MAMMA1001969	F-MAMMA1001969	0.31
15	MAMMA1001970	F-MAMMA1001970	0.34
	MAMMA1002009	F-MAMMA1002009	0.45
	MAMMA1002011	F-MAMMA1002011,	0.94
	MAMMA1002033	F-MAMMA1002033	0.33
20	MAMMA1002041	F-MAMMA1002041	0.86
	MAMMA1002042	F-MAMMA1002042	0.88
	MAMMA1002047	F-MAMMA1002047	0.41
	MAMMA1002068	F-MAMMA1002068	0.38
25	MAMMA1002143	F-MAMMA1002143	0.91
	MAMMA1002153	F-MAMMA1002153	0.94
	MAMMA1002156	F-MAMMA1002156	0.73
	MAMMA1002170	F-MAMMA1002170	0.43
30	MAMMA1002174	F-MAMMA1002174	0.69
	MAMMA1002198	F-MAMMA1002198	0.65
	MAMMA1002209	F-MAMMA1002209	0.80
	MAMMA1C02219	F-MAMMA1002219	0.85
35	MAMMA1002236	F-MAMMA1002236	0.55
	MAMMA1002243	F-MAMMA1002243	0.50
	MAMMA1002268	F-MAMMA1002268	0.43
	MAMMA1002269	F-MAMMA1002269	0.33
40	MAMMA1002292	F-MAMMA1002292	0.76
40	MAMMA1002294	F-MAMMA1002294	0.94
	MAMMA1002297	F-MAMMA1002297	0.91
	MAMMA1002308.	F-MAMMA1002308	0.68
	MAMMA1002312	F-MAMMA1002312	0.77
45	MAMMA1002317	F-MAMMA1002317	0.46
	MAMMA1002319	F-MAMMA1002319	0.31
	MAMMA1002329 MAMMA1002333	F-MAMMA1002329	0.34
	MAMMA1002333	F-MAMMA1002333	0.65
50	MAMMA1002351	F-MAMMA1002351 F-MAMMA1002353	0. 89 0. 94
	MAMMA1002355	F-MAMMA1002355	0. 55
υ	MAMMA1002356	F-MAMMA1002356	0. 33
	MAMMA1002362	F-MAMMA1002362	0. 32
55	MAMMA1002382	F-MAMMA1002382	0. 94
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	MAMMA1002384	F-MAMMA1002384	0. 42
	MAMMA1002385	F-MAMMA1002385	0. 82
	MAMMA1002413	F-MAMMA1002413	0.94
5	MAMMA1002427	F-MAMMA1002427	0.40
	MAMMA1002434	F-MAMMA1002434	0. 74
	MAMMA1002454	F-MAMMA1002454	0. 82
	MAMMA1002461	F-MAMMA1002461	0.49
10	MAMMA1002470	F-MAMMA1002470	0.89
	MAMMA1002485	F-MAMMA1002485	0. 55
	MAMMA1002524	F-MAMMA1002524	0. 33
	MAMMA1002530	F-MAMMA1002530	0.75
15	MAMMA1002554	F-MAMMA1002554	0.41
	MAMMA1002585	F-MAMMA1002585	0.31
	MAMMA1002590	F-MAMMA1002590,	0.72
	MAMMA1002598	F-MAMMA1002598	0.76
20	MAMMA1002617	F-MAMMA1002617	0.49
	MAMMA1002619	F-MAMMA1002619	0.88
	MAMMA1002637	F-MAMMA1002637	0.94
	MAMMA1002650	F-MAMMA1002650	0. 54
25	MAMMA1002655	F-MAMMA1002655	0.39
	MAMMA1002665	F-MAMMA1002665	0.33
	MAMMA1002671	F-MAMMA1002671	0.94
	MAMMA1002673	F-MAMMA1002673	0.70
30	MAMMA1002684	F-MAMMA1002684	0. 94
	MAMMA1002685	F-MAMMA1002685	0.88
	MAMMA1002699	F-MAMMA1002699	0.94
	MAMMA1002711	F-MAMMA1002711	0.39
35	MAMMA1002769	F-MAMMA1002769	0. 88
	MAMMA1002775	F-MAMMA1002775	0. 81
	MAMMA1002782	F-MAMMA1002782	0. 87
	MAMMA1002796	F-MAMMA1002796	0. 40
40	MAMMA1002807	F-MAMMA1002807	0. 64
	MAMMA1002842	F-MAMMA1002842	0. 94
	MAMMA1002843	F-MAMMA1002843	0.32
	MAMMA1002868 MAMMA1002869	F-MAMMA1002868	0. 54
45		F-MAMMA1002869	0.94
45	MAMMA1002881	F-MAMMA1002881	0.94
	MAMMA1002886 MAMMA1002890	F-MAMMA1002886 F-MAMMA1002890	0.61
			0. 94
	MAMMA1002895 MAMMA1002937	F-MAMMA1002895 F-MAMMA1002937	0.94
50	MAMMA1002937	F-MAMMA1002938	0. 91 0. 78
	MAMMA1002938	F-MAMMA1002947	0.78
	MAMMA1002964	F-MAMMA1002964	0. 34
	MAMMA1002987	F-MAMMA1002987	0. 74
55	MAMMA1003011	F-MAMMA1003011	0. 90
			0. 50

	MAMMA1003013	F-MAMMA1003013	0.36
	MAMMA1003015	F-MAMMA1003015	0. 92
_	MAMMA1003035	F-MAMMA1003035	0.94
5	MAMMA1003044	F-MAMMA1003044	0.61
	MAMMA1003047	F-MAMMA1003047	0.94
	MAMMA1003049	F-MAMMA1003049	0.31
	MAMMA1003056	F-MAMMA1003056	0.75
10	MAMMA1003057	F-MAMMA1003057	0.33
	MAMMA1003066	F-MAMMA1003066	0.38
	MAMMA1003099	F-MAMMA1003099	0.79
	MAMMA1003104	F-MAMMA1003104	0.44
15	MAMMA1003113	F-MAMMA1003113	0.90
	MAMMA1003127	F-MAMMA1003127	0.53
	MAMMA1003135	F-MAMMA1003135,	0.94
	MAMMA1003146	F-MAMMA1003146	0.88
20	MAMMA1003150	F-MAMMA1003150	0.73
	MAMMA1003166	F-MAMMA1003166	0.94
	NT2RM1000001	F-NT2RM1000001	0.66
	NT2RM1000018	F-NT2RM1000018	0.55
25	NT2RM1000032	F-NT2RM1000032	0. 74
	NT2RM1000035	F-NT2RM1000035	0. 52
	NT2RM1000037	F-NT2RM1000037	0. 94
	NT2RM1000039	F-NT2RM1000039	0. 77
30	NT2RM1000055	F-NT2RM1000055	0.89
	NT2RM1000059	F-NT2RM1000059	0. 41
	NT2RM1000062	F-NT2RM1000062	0.65
	NT2RM1000080	F-NT2RM1000080	0. 61
35	NT2RM1000086	F-NT2RM1000086	0. 94
	NT2RM1000092	F-NT2RM1000092	0. 94
	NT2RM1000118	F-NT2RM1000118	0.76
	NT2RM1000119	F-NT2RM1000119	0.41
40	NT2RM1000127	F-NT2RM1000127	0. 94
40	NT2RM1000131	F-NT2RM1000131	0.90
	NT2RM1000132	F-NT2RM1000132	0. 94
	NT2RM1000153	F-NT2RM1000153	0. 90
	NT2RM1000186	F-NT2RM1000186	0. 78
45	NT2RM1000187	F-NT2RM1000187	
	NT2RM1000199	F-NT2RM1000199	0. 33
	NT2RM1000242	F-NT2RM1000242	0. 78
	NT2RM1000244	F-NT2RM1000244	0. 31
50	NT2RM1000252	F-NT2RM1000252	0.76
	NT2RM1000256	F-NT2RM1000256	0. 94
	NT2RM1000257	F-NT2RM1000257	0.89
	NT2RM1000260	F-NT2RM1000260	0. 42
55	NT2RM1000271	F-NT2RM1000271	0. 74
	NT2RM1000272	F-NT2RM1000272	0. 78

	NT2RM1000280	F-NT2RM1000280	0. 94
	NT2RM1000300	F-NT2RM1000300	0. 47
	NT2RM1000314	F-NT2RM1000314	0.61
5	NT2RM1000318	F-NT2RM1000318	0.94
	NT2RM1000341	F-NT2RM1000341	0.70
	NT2RM1000354	F-NT2RM1000354	0. 82
	NT2RM1000355	F-NT2RM1000355	0. 94
10	NT2RM1000365	F-NT2RM1000365	0. 31
	NT2RM1000377	F-NT2RM1000377	0.87
	NT2RM1000377	F-NT2RM1000388	0. 94
	NT2RM1000394	F-NT2RM1000394	0. 58
15	NT2RM1000399	F-NT2RM1000399	0.86
73	NT2RM1000421	F-NT2RM1000421	0.39
	NT2RM1000421	F-NT2RM1000430,	
	NT2RM1000499	F-NT2RM1000499	0.44
	NT2RM1000539	F-NT2RM1000539	0.87
20	NT2RM1000553	F-NT2RM1000553	0.88
	NT2RM1000555	F-NT2RM1000555	0. 37
	NT2RM1000563	F-NT2RM1000563	0.63
	NT2RM1000623	F-NT2RM1000623	0.34
25	NT2RM1000648	F-NT2RM1000648	0.94
	NT2RM1000661	F-NT2RM1000661	0.61
	NT2RM1000666	F-NT2RM1000666	0.94
	NT2RM1000669	F-NT2RM1000669	0.94
30	NT2RM1000672	F-NT2RM1000672	0.53
	NT2RM1000691	F-NT2RM1000691	0.38
	NT2RM1000699	F-NT2RM1000699	0.32
•	NT2RM1000702	F-NT2RM1000702	0.74
35	NT2RM1000725	F-NT2RM1000725	0.75
	NT2RM1000741	F-NT2RM1000741	0.94
	NT2RM1000742	F-NT2RM1000742	0.54
	NT2RM1000746	F-NT2RM1000746	0.94
40	NT2RM1000770	F-NT2RM1000770	0.48
	NT2RM1000772	F-NT2RM1000772	0.39
	NT2RM1000780	F-NT2RM1000780	0.66
	NT2RM1000781	F-NT2RM1000781	0.49
45	NT2RM1000800	F-NT2RM1000800	0.72
	NT2RM1000802	F-NT2RM1000802	0.89
	NT2RM1000811	F-NT2RM1000811	0.53
	NT2RM1000826	F-NT2RM1000826	0.39
50	NT2RM1000829	F-NT2RM1000829	0. 63
	NT2RM1000833	F-NT2RM1000833	0. 94
	NT2RM1000850	F-NT2RM1000850	0.89
	NT2RM1000852	F-NT2RM1000852	0.88
56	NT2RM1000857	F-NT2RM1000857	0.58
55	NT2RM1000867	F-NT2RM1000867	0.90

	NT2RM1000874	F-NT2RM1000874	0. 89
	NT2RM1000882	F-NT2RM1000882	0. 63
	NT2RM1000883	F-NT2RM1000883	0. 71
5	NT2RM1000885	F-NT2RM1000885	0. 89
	NT2RM1000894	F-NT2RM1000894	0. 57
	NT2RM1000898	F-NT2RM1000898	0. 87
	NT2RM1000905	F-NT2RM1000905	0. 94
10	NT2RM1000924	F-NT2RM1000924	0. 88
	NT2RM1000927	F-NT2RM1000927	0. 70
	NT2RM1000962	F-NT2RM1000962	0. 84
	NT2RM1000978	F-NT2RM1000978	0. 39
15	NT2RM1000978	F-NT2RM1000378	0. 81
	NT2RM1001008	F-NT2RM1001008	0. 94
	NT2RM1001003	F-NT2RM1001043,	
	NT2RM1001044	F-NT2RM1001044	0. 94
20	NT2RM1001044	F-NT2RM1001059	0. 94
20	NT2RM1001055	F-NT2RM1001066	0. 94
	NT2RM1001072	F-NT2RM1001072	0. 77
	NT2RM1001074	F-NT2RM1001074	0. 94
	NT2RM1001082	F-NT2RM1001082	0. 94
25	NT2RM1001085	F-NT2RM1001085	0. 83
	NT2RM1001092	F-NT2RM1001092	0.77
	NT2RM1001102	F-NT2RM1001102	0. 59
	NT2RM1001105	F-NT2RM1001105	0.94
30	NT2RM1001112	F-NT2RM1001112	0.50
	NT2RM1001115	F-NT2RM1001115	0. 32
	NT2RM1001139	F-NT2RM1001139	0. 77
	NT2RM2000006	F-NT2RM2000006	0.34
35	NT2RM2000013	F-NT2RM2000013	0.70
	NT2RM2000030	F-NT2RM2000030	0.89
	NT2RM2000032	F-NT2RM2000032	0.36
	NT2RM2000042	F-NT2RM2000042	0.32
40	NT2RM2000092	F-NT2RM2000092	0.55
	NT2RM2000093	F-NT2RM2000093	0. 33
	NT2RM2000101	F-NT2RM2000101	0.77
	NT2RM2000124	F-NT2RM2000124	0. 94
45	NT2RM2000191	F-NT2RM2000191	0.80
	NT2RM2000192	F-NT2RM2000192	0.93
	NT2RM2000239	F-NT2RM2000239	0.68
	NNNNNNNNNNN	F-NNNNNNNNNNNN	0.94
50	NT2RM2000250	F-NT2RM2000250	0.94
	NT2RM2000259	F-NT2RM2000259	0.68
	NT2RM2000260	F-NT2RM2000260	0.37
	NT2RM2000287	F-NT2RM2000287	0.46
55	NT2RM2000322	F-NT2RM2000322	0.83
<del></del>	NT2RM2000359	F-NT2RM2000359	0.94

	NT2RM2000363	F-NT2RM2000363	0. 91
	NT2RM2000368	F-NT2RM2000368	0. 34
	NT2RM2000371	F-NT2RM2000371	0. 94
5	NT2RM2000374	F-NT2RM2000374_	
	NT2RM2000395	F-NT2RM20003/4_	0.33
	NT2RM2000393	F-NT2RM2000402	0. 80
10	NT2RM2000407	F-NT2RM2000407	0.88
	NT2RM2000420	F-NT2RM2000420	0.92
	NT2RM2000422	F-NT2RM2000422	0.49
	NT2RM2000452	F-NT2RM2000452	0. 93
15	NT2RM2000469	F-NT2RM2000469	0.39
-	NT2RM2000490	F-NT2RM2000490	0.90
	NT2RM2000502	F-NT2RM2000502	0.64
	NT2RM2000504	F-NT2RM2000504,	0.71
20	NT2RM2000522	F-NT2RM2000522	0.91
20	NT2RM2000540	F-NT2RM2000540	0.87
	NT2RM2000556		0.91
	NT2RM2000566	F-NT2RM2000566	0.39
	NT2RM2000567	F-NT2RM2000567	0. 44
25	NT2RM2000569	F-NT2RM2000569	0.49
	NT2RM2000577	F-NT2RM2000577	0. 45
	NT2RM2000581	F-NT2RM2000581	0. 72
	NT2RM2000588	F-NT2RM2000588	0.31
30	NT2RM2000594	F-NT2RM2000594 F-NT2RM2000599	0.79
	NT2RM2000599 NT2RM2000612	F-NT2RM2000612	0. 80 0. 75
	NT2RM2000612	F-NT2RM2000612	0. 75
	NT2RM2000623	F-NT2RM2000624	0. 72
35	NT2RM2000624	F-NT2RM2000635	0. 73
	NT2RM2000636	F-NT2RM2000636	0. 75
	NT2RM2000639	F-NT2RM2000639	0. 43
	NT2RM2000649	F-NT2RM2000649	0. 74
40	NT2RM2000669	F-NT2RM2000669	0. 94
	NT2RM2000691	F-NT2RM2000691	0.84
	NT2RM2000714	F-NT2RM2000714	0. 92
	NT2RM2000718	F-NT2RM2000718	0. 52
45	NT2RM2000735	F-NT2RM2000735	0. 33
	NT2RM2000740	F-NT2RM2000740	0. 94
	NT2RM2000795	F-NT2RM2000795	0. 44
	NT2RM2000821	F-NT2RM2000821	0. 33
50	NT2RM2000837	F-NT2RM2000837	0. 57
	NT2RM2000951	F-NT2RM2000951	0. 32
	NT2RM2000952	F-NT2RM2000952	0.50
	NT2RM2000984	F-NT2RM2000984	0. 76
55	NT2RM2001004	F-NT2RM2001004	0.55
	NT2RM2001035	F-NT2RM2001035	0. 82

	NT2RM2001065	F-NT2RM2001065	0. 65
	NT2RM2001100	F-NT2RM2001100	0.89
	NT2RM2001105	F-NT2RM2001105	0. 94
5	NT2RM2001103	F-NT2RM2001103	
		F-NT2RM2001131_	0.74
	NT2RM2001141	F-NT2RM2001141	
	NT2RM2001152		0. 33
10	NT2RM2001177	F-NT2RM2001177	0. 32
	NT2RM2001194	F-NT2RM2001194	0. 83
	NT2RM2001196	F-NT2RM2001196	0. 42
	NT2RM2001201	F-NT2RM2001201	0. 90
15	NT2RM2001221	F-NT2RM2001221	0.39
15	NT2RM2001238	F-NT2RM2001238	0. 94
	NT2RM2001243	F-NT2RM2001243	0. 44
	NT2RM2001247	F-NT2RM2001247	
	NT2RM2001256	F-NT2RM2001256	0.60
20	NT2RM2001291	F-NT2RM2001291	0. 39
	NT2RM2001306	F-NT2RM2001306	0. 81
	NT2RM2001312	F-NT2RM2001312	0. 64
	NT2RM2001319	F-NT2RM2001319	0. 53
25	NT2RM2001324	F-NT2RM2001324	0. 94
	NT2RM2001345	F-NT2RM2001345	0. 94
	NT2RM2001360	F-NT2RM2001360	0.69
	NT2RM2001370	F-NT2RM2001370	0. 44
30	NT2RM2001393	F-NT2RM2001393	0. 94
	NT2RM2001420	F-NT2RM2001420	0. 46
	NT2RM2001424	F-NT2RM2001424	0. 94
	NT2RM2001499	F-NT2RM2001499	0. 63
35	NT2RM2001504	F-NT2RM2001504	0. 94
	NT2RM2001524	F-NT2RM2001524	0. 76
,	NT2RM2001544	F-NT2RM2001544	0. 59
	NT2RM2001547	F-NT2RM2001547	0. 48
	NT2RM2001575	F-NT2RM2001575	0. 91
40	NT2RM2001582	F-NT2RM2001582	0. 34
	NT2RM2001588	F-NT2RM2001588	0. 48
	NT2RM2001592	F-NT2RM2001592	0.64
	NT2RM2001605	F-NT2RM2001605	0.47
45	NT2RM2001613	F-NT2RM2001613	0. 61
	NT2RM2001632	F-NT2RM2001632	0. 31
	NT2RM2001635	F-NT2RM2001635	0. 39
	NT2RM2001637	F-NT2RM2001637	0.71
50	NT2RM2001641	F-NT2RM2001641	0. 73
	NT2RM2001648	F-NT2RM2001648	0.44
	NT2RM2001652	F-NT2RM2001652	0.40
	NT2RM2001659	F-NT2RM2001659	0.35
55	NT2RM2001664	F-NT2RM2001664	0.53
	NT2RM2001668	F-NT2RM2001668	0.64

	NT2RM2001670 F-NT2RM2001670	0.59
	NT2RM2001671 F-NT2RM2001671	0.41
	NT2RM2001675 F-NT2RM2001675	0.51
5	NT2RM2001681 F-NT2RM2001681_	0.33
	NT2RM2001688 F-NT2RM2001688	0.74
	NT2RM2001695 F-NT2RM2001695	0.31
	NT2RM2001696 F-NT2RM2001696	0.94
10	NT2RM2001698 F-NT2RM2001698	0.69
	NT2RM2001699 F-NT2RM2001699	0.85
	NT2RM2001700 F-NT2RM2001700	.0. 71
	NT2RM2001706 F-NT2RM2001706	0.38
15	NT2RM2001716 F-NT2RM2001716	0.78
	NT2RM2001718 F-NT2RM2001718	0.81
	NT2RM2001723 F-NT2RM2001723,	0.83
	NT2RM2001727 F-NT2RM2001727	0.87
20	NT2RM2001730 F-NT2RM2001730	0.94
20	NT2RM2001743 F-NT2RM2001743	0.94
	NT2RM2001753 F-NT2RM2001753	0.75
	NT2RM2001760 F-NT2RM2001760	0.75
25	NT2RM2001768 F-NT2RM2001768	0.71
25	NT2RM2001771 F-NT2RM2001771	0.33
	NT2RM2001782 F-NT2RM2001782	0.34
	NT2RM2001784 F-NT2RM2001784	0.45
	NT2RM2001785 F-NT2RM2001785	0.94
30	NT2RM2001797 F-NT2RM2001797	0.74
	NT2RM2001800 F-NT2RM2001800	0.90
	NT2RM2001803 F-NT2RM2001803	0.36
	NT2RM2001805 F-NT2RM2001805	0. 81
35	NT2RM2001813 F-NT2RM2001813	0. 94
	NT2RM2001823 F-NT2RM2001823	0. 94
	NT2RM2001839 F-NT2RM2001839	0. 39
	NT2RM2001840 F-NT2RM2001840	0. 94
40	NT2RM2001855 F-NT2RM2001855	0.92
	NT2RM2001867 F-NT2RM2001867	0.82
	NT2RM2001879 F-NT2RM2001879	0.86
	NT2RM2001886 F-NT2RM2001886	0.91
45	NT2RM2001896 F-NT2RM2001896	0.94
	NT2RM2001903 F-NT2RM2001903	0.61
	NT2RM2001930 F-NT2RM2001930	0.94
	NT2RM2001935 F-NT2RM2001935	0.77
50	NT2RM2001936 F-NT2RM2001936	0.94
	NT2RM2001950 F-NT2RM2001950	0. 34
	NT2RM2001982 F-NT2RM2001982	0.48
	NT2RM2001983 F-NT2RM2001983	0.59
55	NT2RM2001989 F-NT2RM2001989	0.94
	NT2RM2001997 F-NT2RM2001997	0.90

	NT2RM2001998	F-NT2RM2001998	0. 72
	NT2RM2002004	F-NT2RM2002004	0.46
	NT2RM2002014	F-NT2RM2002014	0. 32
5	NT2RM2002030	F-NT2RM2002030_	0.54
	NT2RM2002049	F-NT2RM2002049	0. 59
	NT2RM2002055	F-NT2RM2002055	0.94
	NT2RM2002088	F-NT2RM2002088	0.47
10	NT2RM2002091	F-NT2RM2002091	0. 78-
•	NT2RM2002100	F-NT2RM2002100	0. 87
	NT2RM2002109	F-NT2RM2002109	0.86
	NT2RM2002128	F-NT2RM2002128	0.42
15	NT2RM2002142	F-NT2RM2002142	0. 94
	NT2RM2002145	F-NT2RM2002145	0.94
	NT2RM2002178	F-NT2RM2002178;	0.44
	NT2RM2002580	F-NT2RM2002580	0.94
20	NT2RM4000024	F-NT2RM4000024	0.76
	NT2RM4000027	F-NT2RM4000027	0.40
	NT2RM4000030	F-NT2RM4000030	0.66
	NT2RM4000061	F-NT2RM4000061	0. 33
25	NT2RM4000104	F-NT2RM4000104	0.93
	NT2RM4000139	F-NT2RM4000139	0.87
	NT2RM4000155	F-NT2RM4000155	0.51
	NT2RM4000156	F-NT2RM4000156	0.39
30	NT2RM4000167	F-NT2RM4000167	0.50
	NT2RM4000169	F-NT2RM4000169	0.44
	NT2RM4000191	F-NT2RM4000191	0.94
	NT2RM4000197	F-NT2RM4000197	0.41
35	NT2RM4000199	F-NT2RM4000199	0.76
	NT2RM4000202	F-NT2RM4000202	0. 47
	NT2RM4000210	F-NT2RM4000210	0.44
	NT2RM4000215	F-NT2RM4000215	0. 94
	NT2RM4000229	F-NT2RM4000229	0. 93
40	NT2RM4000233	F-NT2RM4000233	0. 74
	NT2RM4000251	F-NT2RM4000251	0. 48
	NT2RM4000290	F-NT2RM4000290	0. 94
-	NT2RM4000324	F-NT2RM4000324	0. 32
45	NT2RM4000344	F-NT2RM4000344	0. 94
·	NT2RM4000349	F-NT2RM4000349	0. 78
	NT2RM4000354	F-NT2RM4000354	0. 36
	NT2RM4000356	F-NT2RM4000356	0.58
50	NT2RM4000386	F-NT2RM4000386	0. 90
	NT2RM4000395	F-NT2RM4000395	0.47
	NT2RM4000421	F-NT2RM4000421	0.46
	NT2RM4000433	F-NT2RM4000433	0.48
55	NT2RM4000457	F-NT2RM4000457	0.75
	NT2RM4000471	F-NT2RM4000471	0. 92

	NT2RM4000486	F-NT2RM4000486	0. 55
	NT2RM4000496	F-NT2RM4000496	0. 62
	NT2RM4000511	F-NT2RM4000511	0.74
5	NT2RM4000514	F-NT2RM4000514	0.72
	NT2RM4000515	F-NT2RM4000515	0.66
	NT2RM4000520	F-NT2RM4000520	0.69
	NT2RM4000531	F-NT2RM4000531	0, 62
10	NT2RM4000534	F-NT2RM4000534	0. 94
	NT2RM4000585	F-NT2RM4000585	0. 52
	NT2RM4000590	F-NT2RM4000590	0.94
	NT2RM4000595	F-NT2RM4000595	0.44
15	NT2RM4000603	F-NT2RM4000603	0.94
	NT2RM4000611	F-NT2RM4000611	0.46
	NT2RM4000616	F-NT2RM4000616,	
	NT2RM4000674	F-NT2RM4000674	0.80
20	NT2RM4000689	F-NT2RM4000689	0.58
	NT2RM4000698	F-NT2RM4000698	0.71
	NT2RM4000700	F-NT2RM4000700	0. 59
	NT2RM4000712	F-NT2RM4000712	0.94
25	NT2RM4000717	F-NT2RM4000717	0.69
	NT2RM4000733	F-NT2RM4000733	0.94
•	NT2RM4000734	F-NT2RM4000734	0.42
	NT2RM4000741	F-NT2RM4000741	0.83
30	NT2RM4000751	F-NT2RM4000751	0.91
	NT2RM4000764	F-NT2RM4000764	0.35
	NT2RM4000778	F-NT2RM4000778	0.90
	NT2RM4000787	F-NT2RM4000787	0.40
95	NT2RM4000790	F-NT2RM4000790	0.62
35	NT2RM4000795	F-NT2RM4000795	0.94
	NT2RM4000796	F-NT2RM4000796	0. 93
	NT2RM4000798	F-NT2RM4000798	0.61
	NT2RM4000813	F-NT2RM4000813	0.58
40	NT2RM4000820	F-NT2RM4000820	0.39
	NT2RM4000833	F-NT2RM4000833	0.80
	NT2RM4000848	F-NT2RM4000848	0.73
	NT2RM4000852	F-NT2RM4000852	0.74
45	NT2RM4000887	F-NT2RM4000887	0.42
	NT2RM4000895	F-NT2RM4000895	0.47
	NT2RM4000950	F-NT2RM4000950	0.53
	NT2RM4000971	F-NT2RM4000971	0.73
50	NT2RM4000996	F-NT2RM4000996	0.42
	NT2RM4001002	F-NT2RM4001002	0.57
	NT2RM4001032	F-NT2RM4001032	0.94
	NT2RM4001047	F-NT2RM4001047	0. 76
55	NT2RM4001054	F-NT2RM4001054	0.91
	NT2RM4001084	F-NT2RM4001084	0. 94

	NT2RM4001092 F-NT2RM4001092 (	). 94
	NT2RM4001116 F-NT2RM4001116 (	0. 36
	NT2RM4001140 F-NT2RM4001140 (	0. 82
5	NT2RM4001151 F-NT2RM4001151(	0.94
		0.94
		0. 61
		0.77
10	WI ZIMI 100 TIO	0.72
10	11/2/(11/100/200	0.60
	11/2/11/1007	0.89
	1412///1400120	0. 93
	****	0. 89
15	WIZIGIFTOOTZOS	0. 94
	14121/1117001200	0. 88
	RTZIMITOOTOO , III ZIIII	0. 38
	111210111001010	0. 33
20	W1210H7001020	0. 33 .0, 83
	MIZAMI POST A PARTICIPANT A PA	0. 94
	11/2/10/10/10/1	0.88
	161211111101011	0.61
25	NT2RM4001371 F-NT2RM4001371	0. 93
	NT2RM4001382 F-NT2RM4001382	
	NT2RM4001384 F-NT2RM4001384	0.32
	NT2RM4001410 F-NT2RM4001410	0.94
20	NT2RM4001411 F-NT2RM4001411	0.47
30	NT2RM4001412 F-NT2RM4001412	0.85
	NT2RM4001444 F-NT2RM4001444	0.94
•	NT2RM4001454 F-NT2RM4001454	0.72
	NT2RM4001455 F-NT2RM4001455	0.39
35	NT2RM4001483 F-NT2RM4001483	0.76
	NT2RM4001489 F-NT2RM4001489	0.94
	NT2RM4001522 F-NT2RM4001522	0.88
	NT2RM4001565 F-NT2RM4001565	0.78
40	NT2RM4001566 F-NT2RM4001566	0.87
	NT2RM4001569 F-NT2RM4001569	0.70
	NT2RM4001582 F-NT2RM4001582	0.76
	NT2RM4001592 F-NT2RM4001592	0.31
45	NT2RM4001594 F-NT2RM4001594	0.86
	NT2RM4001597 F-NT2RM4001597	0.91
	NT2RM4001611 F-NT2RM4001611	0.91
	NT2RM4001629 F-NT2RM4001629	0.72
50	NT2RM4001650 F-NT2RM4001650	0.37
50	NT2RM4001662 F-NT2RM4001662	0.70
	NT2RM4001666 F-NT2RM4001666	0.37
	NT2RM4001682 F-NT2RM4001682	0.91
	NT2RM4001710 F-NT2RM4001710	0. 45
55	NT2RM4001714 F-NT2RM4001714	0. 94
	•	

	NT2RM4001715	F-NT2RM4001715	0. 33
	NT2RM4001731	F-NT2RM4001731	0.94
	NT2RM4001741	F-NT2RM4001741	0.87
5	NT2RM4001746	F-NT2RM4001746	0.81
	NT2RM4001758	F-NT2RM4001758	0.59
	NT2RM4001783	F-NT2RM4001783	0.68
	NT2RM4001813	F-NT2RM4001813	0.39
10	NT2RM4001819	F-NT2RM4001819	0. 43
	NT2RM4001823	F-NT2RM4001823	0. 94
	NT2RM4001828	F-NT2RM4001828	0. 83
	NT2RM4001836	F-NT2RM4001836	0. 61
15	NT2RM4001841	F-NT2RM4001841	0.37
	NT2RM4001842	F-NT2RM4001842	0. 68
	NT2RM4001865	F-NT2RM4001865	
	NT2RM4001876	F-NT2RM4001876	0. 88
	NT2RM4001880	F-NT2RM4001880	0. 43
20	NT2RM4001905	F-NT2RM4001905	0. 94
	NT2RM4001922	F-NT2RM4001922	0. 48
	NT2RM4001938	F-NT2RM4001938	0. 67
	NT2RM4001940	F-NT2RM4001940	0. 92
25	NT2RM4001965	F-NT2RM4001965	0. 94
	NT2RM4001969	F-NT2RM4001969	0. 88
	NT2RM4001979	F-NT2RM4001979	0. 42
	NT2RM4001987	F-NT2RM4001987	0. 86
30	NT2RM4002013	F-NT2RM4002013	0. 94
	NT2RM4002018	F-NT2RM4002018	0. 92
	NT2RM4002034	F-NT2RM4002034	0. 43
	NT2RM4002044	F-NT2RM4002044	0. 42
35	NT2RM4002054	F-NT2RM4002054	0.88
	NT2RM4002055	F-NT2RM4002055	0. 32
	NT2RM4002062	F-NT2RM4002062	0. 87
	NT2RM4002063	F-NT2RM4002063	0. 49
40	NT2RM4002066	F-NT2RM4002066	0. 39
40	NT2RM4002073	F-NT2RM4002073	0. 78
	NT2RM4002075	F-NT2RM4002075	0. 87
	NT2RM4002093	F-NT2RM4002093	0. 82
	NT2RM40C2109	F-NT2RM4002109	0. 52
45	NT2RM4002128	F-NT2RM4002128	0. 68
	NT2RM4002140	F-NT2RM4002140	0. 94
	NT2RM4002145	F-NT2RM4002145	0. 94
	NT2RM4002146	F-NT2RM4002146	0. 89
50	NT2RM4002140	F-NT2RM4002161	0. 49
	NT2RM4002174	F-NT2RM4002174	0. 36
	NT2RM4002174	F-NT2RM4002174	0. 34
	NT2RM4002194	F-NT2RM4002194	0. 73
55	NT2RM4002194	F-NT2RM4002205	0. 94
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	NT2RM4002213	F-NT2RM4002213	0.52
	NT2RM4002226	F-NT2RM4002226	0.48
	NT2RM4002251	F-NT2RM4002251	0.94
5	NT2RM4002256	F-NT2RM4002256_	0.93
	NT2RM4002266	F-NT2RM4002266	0.73
	NT2RM4002294	F-NT2RM4002294	0.86
	NT2RM4002301	F-NT2RM4002301	0.40
10	NT2RM4002323	F-NT2RM4002323	0. 81
	NT2RM4002339	F-NT2RM4002339	0.32
	NT2RM4002344	F-NT2RM4002344	0.87
	NT2RM4002373	F-NT2RM4002373	0.41
15	NT2RM4002374	F-NT2RM4002374	0.33
	NT2RM4002398	F-NT2RM4002398	0.94
	NT2RM4002409	F-NT2RM4002409	
	NT2RM4002446	F-NT2RM4002446	0. 61
. 20	NT2RM4002452	F-NT2RM4002452	0.47
	NT2RM4002457	F-NT2RM4002457	0.92
	NT2RM4002460	F-NT2RM4002460	0.61
	NT2RM4002482	F-NT2RM4002482	0. 94
25	NT2RM4002493	F-NT2RM4002493	0.39
23	NT2RM4002527	F-NT2RM4002527	0.64
	NT2RM4002534	F-NT2RM4002534	0.76
	NT2RM4002558	F-NT2RM4002558	0.62
00	NT2RM4002565	F-NT2RM4002565	0.54
30	NT2RM4002571	F-NT2RM4002571	0.90
	NT2RM4002593	F-NT2RM4002593	0.91
	NT2RM4002594	F-NT2RM4002594	0.70
	NT2RM4002623	F-NT2RM4002623	0.81
35	NT2RP1000018	F-NT2RP1000018	0.31
	NT2RP1000035	F-NT2RP1000035	0.49
	NT2RP1000040	F-NT2RP1000040	0.45
	NT2RP1000063	F-NT2RP1000063	0.90
40	NT2RP1000086	F-NT2RP1000086	0.89
	NT2RP1000101	F-NT2RP1000101	0.76
	NT2RP1000111	F-NT2RP1000111	0.94
	NT2RP1000112	F-NT2RP1000112	0. 45
45	NT2RP1000124	F-NT2RP1000124	0. 94
	NT2RP1000130	F-NT2RP1000130	0.69
	NT2RP1000163	F-NT2RP1000163	0.90
	NT2RP1000170	F-NT2RP1000170	0.94
50	NT2RP1000174	F-NT2RP1000174	0.74
	NT2RP1000191	F-NT2RP1000191	0.76
	NT2RP1000202	F-NT2RP1000202	0.60
	NT2RP1000243	F-NT2RP1000243	0.81
55	NT2RP1000259	F-NT2RP1000259	0.79
	NT2RP1000272	F-NT2RP1000272	0.94

	NT2RP1000324	F-NT2RP1000324	0. 32
	NT2RP1000326	F-NT2RP1000326	0. 94
	NT2RP1000333	F-NT2RP1000333	0.94
5	NT2RP1000348	F-NT2RP1000348_	0.94
	NT2RP1000357	F-NT2RP1000357	0.87
	NT2RP1000358	F-NT2RP1000358	0.79
	NT2RP1000363	F-NT2RP1000363	0.40
10	NT2RP1000376	F-NT2RP1000376	0.61
	NT2RP1000409	F-NT2RP1000409	0.85
	NT2RP1000413	F-NT2RP1000413	0.42
	NT2RP1000416	F-NT2RP1000416	0.68
15	NT2RP1000418	F-NT2RP1000418	0.88
	NT2RP1000439	F-NT2RP1000439	0.69
	NT2RP1000443	F-NT2RP1000443	0. 59
	NT2RP1000460	F-NT2RP1000460	0.86
20	NT2RP1000470	F-NT2RP1000470	0.56
20	NT2RP1000478	F-NT2RP1000478	0.87
	NT2RP1000481	F-NT2RP1000481	0.58
	NT2RP1000493	F-NT2RP1000493	0.34
	NT2RP1000513	F-NT2RP1000513	0.89
25	NT2RP1000522	F-NT2RP1000522	0.34
	NT2RP1000547	F-NT2RP1000547	0.89
	NT2RP1000574	F-NT2RP1000574	0.74
	NT2RP1000577	F-NT2RP1000577	0.53
30	NT2RP1000581	F-NT2RP1000581	0.66
	NT2RP1000609	F-NT2RP1000609	0.90
	NT2RP1000629	F-NT2RP1000629	0.94
	NT2RP1000630	F-NT2RP1000630	0.93
35	NT2RP1000677	F-NT2RP1000677	0. 94
•	NT2RP1000688	F-NT2RP1000688	0. 94
	NT2RP1000695	F-NT2RP1000695	0.89
	NT2RP1000701	F-NT2RP1000701	0.89
40	NT2RP1000721	F-NT2RP1000721	0.91
	NT2RP1000730	F-NT2RP1000730	0. 94
	NT2RP1000733	F-NT2RP1000733	0.90
	NT2RP1000738	F-NT2RP1000738	0.64
45	NT2RP1000746	F-NT2RP1000746	0. 92
	NT2RP1000767	F-NT2RP1000767	0.82
	NT2RP1000782	F-NT2RP1000782	0. 94
	NT2RP1000796	F-NT2RP1000796	0. 33
50	NT2RP1000825	F-NT2RP1000825	0.74
	NT2RP1000833	F-NT2RP1000833	0.94
	NT2RP1000834	F-NT2RP1000834	0.88
	NT2RP1000836	F-NT2RP1000836	0.59
55	NT2RP1000846	F-NT2RP1000846	0.46
	NT2RP1000851	F-NT2RP1000851	0. 82

	NT2RP1000856	F-NT2RP1000856	0.94
	NT2RP1000860	F-NT2RP1000860	0.60
•	NT2RP1000902	F-NT2RP1000902	0.42
5	NT2RP1000915	F-NT2RP1000915	0.90
	NT2RP1000916	F-NT2RP1000916	0. 67
	NT2RP1000943	F-NT2RP1000943	0.77
	NT2RP1000944	F-NT2RP1000944	0.94
10	NT2RP1000947	F-NT2RP1000947	0.38
	NT2RP1000954	F-NT2RP1000954	0.59
	NT2RP1000958	F-NT2RP1000958	0.59
	NT2RP1000959	F-NT2RP1000959	0.74
15	NT2RP1000966	F-NT2RP1000966	0.93
	NT2RP1000980	F-NT2RP1000980	0.31
	NT2RP1000988	F-NT2RP1000988	
	NT2RP1001011	F-NT2RP1001011	0.65
20	NT2RP1001013	F-NT2RP1001013	0.34
	NT2RP1001014	F-NT2RP1001014	0.94
	NT2RP1001033	F-NT2RP1001033	0.40
	NT2RP1001073	F-NT2RP1001073	0.74
05	NT2RP1001079	F-NT2RP1001079	0.48
25	NT2RP1001080	F-NT2RP1001080	0.91
	NT2RP1001113	F-NT2RP1001113	0.94
	NT2RP1001173	F-NT2RP1001173	0.57
	NT2RP1001177	F-NT2RP1001177	0.75
30	NT2RP1001185	F-NT2RP1001185	0.94
	NT2RP1001199	F-NT2RP1001199	0.42
	NT2RP1001247	F-NT2RP1001247	0.67
	NT2RP1001248	F-NT2RP1001248	0.89
35	NT2RP1001253	F-NT2RP1001253	0.94
	NT2RP1001286	F-NT2RP1001286	0.59
	NT2RP1001294	F-NT2RP1001294	0.81
	NT2RP1001302	F-NT2RP1001302	0.86
40	NT2RP1001310	F-NT2RP1001310	0.91
	NT2RP1001311	F-NT2RP1001311	0.87
	NT2RP1001313	F-NT2RP1001313	0.91
	NT2RP1001361	F-NT2RP1001361	0.83
45	NT2RP1001385	F-NT2RP1001385	0.36
	NT2RP1001395	F-NT2RP1001395	0.84
	NT2RP1001410	F-NT2RP1001410	0.72
	NT2RP1001424	F-NT2RP1001424	0.81
50	NT2RP1001432		0.81
	NT2RP1001449	F-NT2RP1001449	0.77
	NT2RP1001457		0.92
	NT2RP1001466		
55	NT2RP1001475		
	NT2RP1001482	F-NT2RP1001482	0.34

	NT2RP1001494	F-NT2RP1001494	0. 93
	NT2RP1001543	F-NT2RP1001543	0. 94
	NT2RP1001546	F-NT2RP1001546	0.94
5	NT2RP1001569	F-NT2RP1001569_	0, 93
	NT2RP1001616	F-NT2RP1001616	0.46
	NT2RP1001665	F-NT2RP1001665	0.94
	NT2RP2000001	F-NT2RP2000001	0.94
10	NT2RP2000006	F-NT2RP2000006	0.74
	NT2RP2000007	F-NT2RP2000007	0.94
	NT2RP2000008	F-NT2RP2000008	0.83
	NT2RP2000032	F-NT2RP2000032	0, 94
15	NT2RP2000040	F-NT2RP2000040	0.92
	NT2RP2000045	F-NT2RP2000045	0.94
	NT2RP2000054	F-NT2RP2000054	0.53
	NT2RP2000056	F-NT2RP2000056	0. 42
20	NT2RP2000067	F-NT2RP2000067	0.88
	NT2RP2000070	F-NT2RP2000070	0.48
	NT2RP2000079	F-NT2RP2000079	0.85
	NT2RP2000088	F-NT2RP2000088	0. 94
25	NT2RP2000091	F-NT2RP2000091	0. 79
	NT2RP2000097	F-NT2RP2000097	0.66
	NT2RP2000114	F-NT2RP2000114	0.86
	NT2RP2000120	F-NT2RP2000120	0.48
30	NT2RP2000126	F-NT2RP2000126	0. 77
	NT2RP2000133	F-NT2RP2000133	0. 64
	NT2RP2000147	F-NT2RP2000147	0. 94
	NT2RP2000153	F-NT2RP2000153	0. 87
35	NT2RP2000157	F-NT2RP2000157	0. 48
	NT2RP2000161	F-NT2RP2000161	0.79
	NT2RP2000173	F-NT2RP2000173	0.75
•	NT2RP2000175	F-NT2RP2000175	0.79
40	NT2RP2000195 NT2RP2000205	F-NT2RP2000195 F-NT2RP2000205	0. 91 0. 94
	NT2RP2000205	F-NT2RP2000208	0. 94
	NT2RP2000208	F-NT2RP2000208	0. 52
•	NT2RP2000224	F-NT2RP2000232	0. 32
45	NT2RP2000232	F-NT2RP2000233	0. 57
	NT2RP2000239	F-NT2RP2000239	0. 52
	NT2RP2000248	F-NT2RP2000248	0.36
	NT2RP2000274	F-NT2RP2000274	0. 55
50	NT2RP2000283	F-NT2RP2000283	0. 61
	NT2RP2000288	F-NT2RP2000288	0. 31
	NT2RP2000297	F-NT2RP2000297	0.78
	NT2RP2000298	F-NT2RP2000298	0, 94
55	NT2RP2000310	F-NT2RP2000310	0. 52
	NT2RP2000328	F-NT2RP2000328	0.75

	NT2RP2000329 F-NT2RP2000329	0. 94
	NT2RP2000346 F-NT2RP2000346	0. 62
	NT2RP2000369 F-NT2RP2000369	0. 87
5	NT2RP2000412 F-NT2RP2000412	0.75
	NT2RP2000414 F-NT2RP2000414	0.79
	NT2RP2000422 F-NT2RP2000422	0.46
	NT2RP2000438 F-NT2RP2000438	0.94
10	NT2RP2000448 F-NT2RP2000448	0.94
	NT2RP2000503 F-NT2RP2000503	0.94
	NT2RP2000510 F-NT2RP2000510	0. 87
	NT2RP2000516 F-NT2RP2000516	0.42
15	NT2RP2000617 F-NT2RP2000617	0.94
	NT2RP2000634 F-NT2RP2000634	0.74
	NT2RP2000656 F-NT2RP2000656,	0. 68
	NT2RP2000658 F-NT2RP2000658	0. 55
20	NT2RP2000668 F-NT2RP2000668	0.81
	NT2RP2000704 F-NT2RP2000704	0.88
	NT2RP2000710 F-NT2RP2000710	0. 58
	NT2RP2000764 F-NT2RP2000764	0.59
25	NT2RP2000809 F-NT2RP2000809	0.51
23	NT2RP2000812 F-NT2RP2000812	0. 37
	NT2RP2000814 F-NT2RP2000814	0.66
	NT2RP2000816 F-NT2RP2000816	0.50
	NT2RP2000819 F-NT2RP2000819	0. 35
30	NT2RP2000841 F-NT2RP2000841	0.93
	NT2RP2000845 F-NT2RP2000845	0.78
	NT2RP2000863 F-NT2RP2000863	0.41
	NT2RP2000880 F-NT2RP2000880	0. 79
35	NT2RP2000892 F-NT2RP2000892	0.94
	NT2RP2000931 F-NT2RP2000931	0. 89
	NT2RP2000932 F-NT2RP2000932	0. 94
	NT2RP2000938 F-NT2RP2000938	0. 90
40	NT2RP2000943 F-NT2RP2000943	0.61
	NT2RP2000965 F-NT2RP2000965	0. 80
	NT2RP2000985 F-NT2RP2000985	0.94
	NT2RP2001036 F-NT2RP2001036	0.86
45	NT2RP2001044 F-NT2RP2001044	0. 93
	NT2RP2001056 F-NT2RP2001056	0. 77
	NT2RP2001065 F-NT2RP2001065	0. 73
	NT2RP2001070 F-NT2RP2001070	•
50	NT2RP2001081 F-NT2RP2001081	0. 94
	NT2RP2001094 F-NT2RP2001094	0. 93
	NT2RP2001119 F-NT2RP2001119	
	NT2RP2001127 F-NT2RP2001127	
55	NT2RP2001137 F-NT2RP2001137	
55	NT2RP2001168 F-NT2RP2001168	0. 71

	NT2RP2001173	F-NT2RP2001173	0. 94
	NT2RP2001174	F-NT2RP2001174	0.71
	NT2RP2001218	F-NT2RP2001218	0.76
5	NT2RP2001233	F-NT2RP2001233	
	NT2RP2001245	F-NT2RP2001245	0.33
	NT2RP2001268	F-NT2RP2001268	0.61
	NT2RP2001327	F-NT2RP2001327	0.86
10	NT2RP2001328	F-NT2RP2001328	0.94
	NT2RP2001366	F-NT2RP2001366	0. 52
	NT2RP2001378	F-NT2RP2001378	0.64
	NT2RP2001381	F-NT2RP2001381	0.80
15	NT2RP2001392	F-NT2RP2001392	0.69
	NT2RP2001394	F-NT2RP2001394	0. 38
	NT2RP2001397	F-NT2RP2001397	
	NT2RP2001420	F-NT2RP2001420	0. 88
20	NT2RP2001427	F-NT2RP2001427	0. 73
20	NT2RP2001436	F-NT2RP2001436	0.32
	NT2RP2001440	F-NT2RP2001440	0.94
	NT2RP2001450	F-NT2RP2001450	0.38
	NT2RP2001511	F-NT2RP2001511	0.53
25	NT2RP2001520	F-NT2RP2001520	0.88
		F-NT2RP2001536	0.64
	NT2RP2001560	F-NT2RP2001560	0.79
	NT2RP2001576	F-NT2RP2001576	0.62
30	NT2RP2001581	F-NT2RP2001581	0.48
	NT2RP2001597	F-NT2RP2001597	0.94
	NT2RP2001601	F-NT2RP2001601	0.33
	NT2RP2001613	F-NT2RP2001613	0.41
35	NT2RP2001628	F-NT2RP2001628	0.79
	NT2RP2001634	F-NT2RP2001634	0.87
	NT2RP2001660	F-NT2RP2001660	0.86
	NT2RP2001675	F-NT2RP2001675	0. 90
40	NT2RP2001721	F-NT2RP2001721	0.32
	NT2RP2001748	F-NT2RP2001748	0.32
	NT2RP2001839	F-NT2RP2001839	0.83
	NT2RP2001861	F-NT2RP2001861	0.33
45	NT2RP2001876	F-NT2RP2001876	0.94
	NT2RP2001883	F-NT2RP2001883	0.90
	NT2RP2001898	F-NT2RP2001898	0.94
	NT2RP2001900	F-NT2RP2001900	0.93
50	NT2RP2001907	F-NT2RP2001907	0.94
	NT2RP2001946	F-NT2RP2001946	0.63
	NT2RP2001947	F-NT2RP2001947	0.75
	NT2RP2001969	F-NT2RP2001969	0.94
55	NT2RP2001976	F-NT2RP2001976	0.48
	NT2RP2001985	F-NT2RP2001985	0. 44

	NT2RP2001991	F-NT2RP2001991	0.39
	NT2RP2002025	F-NT2RP2002025	0.94
	NT2RP2002033	F-NT2RP2002033	0.40
5	NT2RP2002046	F-NT2RP2002046	0.56
	NT2RP2002058	F-NT2RP2002058	0.64
	NT2RP2002076	F-NT2RP2002076	0.40
	NT2RP2002078	F-NT2RP2002078	0.81
10	NT2RP2002079	F-NT2RP2002079	0.82
	NT2RP2002099	F-NT2RP2002099	0.92
	NT2RP2002154	F-NT2RP2002154	0.44
	NT2RP2002185	F-NT2RP2002185	0.94
15	NT2RP2002193	F-NT2RP2002193	0.48
	NT2RP2002208	F-NT2RP2002208	0.49
	NT2RP2002231	F-NT2RP2002231,	0.61
	NT2RP2002235	F-NT2RP2002235	0.87
20	NT2RP2002252	F-NT2RP2002252	0.51
	NT2RP2002270	F-NT2RP2002270	0.78
	NT2RP2002292	F-NT2RP2002292	0.94
	NT2RP2002312	F-NT2RP2002312	0.33
25	NT2RP2002325	F-NT2RP2002325	0.53
	NT2RP2002333	F-NT2RP2002333	0.60
	NT2RP2002373	F-NT2RP2002373	0.91
	NT2RP2002385	F-NT2RP2002385	0.94
30	NT2RP2002408	F-NT2RP2002408	0.44
	NT2RP2002426	F-NT2RP2002426	0.54
•	NT2RP2002442	F-NT2RP2002442	0.37
	NT2RP2002464	F-NT2RP2002464	0.79
35	NT2RP2002479	F-NT2RP2002479	0.92
33	NT2RP2002498	F-NT2RP2002498	0.94
	NT2RP2002503	F-NT2RP2002503	0. 90
	NT2RP2002520	F-NT2RP2002520	0.37
40	NT2RP2002537	F-NT2RP2002537	0.42
40	NT2RP2002549	F-NT2RP2002549	0.75
	NT2RP2002595	F-NT2RP2002595	0.90
	NT2RP2002609	F-NT2RP2002609	0.33
	NT2RP2002618	F-NT2RP2002618	0. 52
45	NT2RP2002621	F-NT2RP2002621	0. 67
	NT2RP2002672	F-NT2RP2002672	0. 43
	NT2RP2002701	F-NT2RP2002701	0.69
	NT2RP2002706	F-NT2RP2002706	0.51
50	NT2RP2002710	F-NT2RP2002710	0.94
	NT2RP2002769	F-NT2RP2002769	0.42
	NT2RP2002800	F-NT2RP2002800	0.47
	NT2RP2002862	F-NT2RP2002862	0. 88 0. 78
55	NT2RP2002880 NT2RP2002891	F-NT2RP2002880 F-NT2RP2002891	0. 78
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	NT2RP2002925	F-NT2RP2002925	0.65
	NT2RP2002928	F-NT2RP2002928	0. 47
	NT2RP2002929	F-NT2RP2002929	0.94
5	NT2RP2002939	F-NT2RP2002939	0.94
	NT2RP2002954	F-NT2RP2002954	0. 94
	NT2RP2002959	F-NT2RP2002959	0. 45
	NT2RP2002979	F-NT2RP2002979	0. 51
10	NT2RP2002980	F-NT2RP2002980	0. 38
	NT2RP2002986	F-NT2RP2002986	0. 34
	NT2RP2002993	F-NT2RP2002993	0.94
	NT2RP2003034	F-NT2RP2003034	0. 34
15	NT2RP2003099	F-NT2RP2003099	0.46
	NT2RP2003108	F-NT2RP2003108	0. 67
	NT2RP2003117	F-NT2RP2003117,	0.57
	NT2RP2003121	F-NT2RP2003121	0.94
20	NT2RP2003125	F-NT2RP2003125	0.94
	NT2RP2003137	F-NT2RP2003137	0.60
	NT2RP2003157	F-NT2RP2003157	0.94
	NT2RP2003158	F-NT2RP2003158	0.75
25	NT2RP2003165	F-NT2RP2003165	0.90
	NT2RP2003177	F-NT2RP2003177	0.85
	NT2RP2003194	F-NT2RP2003194	0.94
	NT2RP2003228	F-NT2RP2003228	0. 57
20	NT2RP2003243	F-NT2RP2003243	0.94
30	NT2RP2003265	F-NT2RP2003265	0.68
	NT2RP2003272	F-NT2RP2003272	0. 94
	NT2RP2003277	F-NT2RP2003277	0. 73
	NT2RP2003280	F-NT2RP2003280	0.94
35	NT2RP2003286	F-NT2RP2003286	0. 44
	NT2RP2003293	F-NT2RP2003293	0.42
	NT2RP2003295	F-NT2RP2003295	0.89
	NT2RP2003297	F-NT2RP2003297	0. 94
40	NT2RP2003307	F-NT2RP2003307	0. 79
	NT2RP2003308	F-NT2RP2003308	0.49
	NT2RP2003329	F-NT2RP2003329	0. 33
	NT2RP2003347	F-NT2RP2003347	0. 79
45	NT2RP2003367	F-NT2RP2003367	0. 94
	NT2RP2003391	F-NT2RP2003391	0.90
	NT2RP2003393	F-NT2RP2003393	0.94
	NT2RP2003401	F-NT2RP2003401	0.80
50	NT2RP2003433	F-NT2RP2003433	0.40
	NT2RP2003445		0. 94 0. 40
	NT2RP2003446 NT2RP2003466		0. 40
	NT2RP2003466 NT2RP2003480		0.58
55			0. 94
	NT2RP2003506	-M  ZR  ZUU3300	0. 94

	NT2RP2003511	F-NT2RP2003511	0: 31
	NT2RP2003513	F-NT2RP2003513	0. 67
	NT2RP2003533	F-NT2RP2003533	0. 57
5	NT2RP2003543	F-NT2RP2003543	0. 94
	NT2RP2003564	F-NT2RP2003564	0. 88
	NT2RP2003567	F-NT2RP2003567	0. 69
	NT2RP2003581	F-NT2RP2003581	0. 40
10	NT2RP2003596	F-NT2RP2003596	0. 83
	NT2RP2003604	F-NT2RP2003604	0. 87
	NT2RP2003629	F-NT2RP2003629	0. 87
	NT2RP2003643	F-NT2RP2003643	0. 94
15	NT2RP2003687	F-NT2RP2003687	0. 31
•	NT2RP2003691	F-NT2RP2003691	0. 34
	NT2RP2003702	F-NT2RP2003702,	
	NT2RP2003713	F-NT2RP2003713	0. 53
20	NT2RP2003714	F-NT2RP2003714	0. 61
	NT2RP2003737	F-NT2RP2003737	0. 88
	NT2RP2003751	F-NT2RP2003751	0.36
	NT2RP2003760	F-NT2RP2003760	0. 62
25	NT2RP2003764	F-NT2RP2003764	0. 94
	NT2RP2003769	F-NT2RP2003769	0. 69
	NT2RP2003777	F-NT2RP2003777	0. 93
	NT2RP2003781	F-NT2RP2003781	0. 37
30	NT2RP2003793	F-NT2RP2003793	0.92
	NT2RP2003825	F-NT2RP2003825	0. 49
	NT2RP2003840	F-NT2RP2003840	0.88
	NT2RP2003857	F-NT2RP2003857	0.34
ar.	NT2RP2003952	F-NT2RP2003952	0.89
35	NT2RP2003976	F-NT2RP2003976	0. 33
	NT2RP2003981	F-NT2RP2003981	0.64
	NT2RP2003984	F-NT2RP2003984	0.41
	NT2RP2003986	F-NT2RP2003986	0. 32
40	NT2RP2004013	F-NT2RP2004013	0.48
	NT2RP2004041	F-NT2RP2004041	0.78
	NT2RP2004042	F-NT2RP2004042	0.66
	NT2RP2004066	F-NT2RP2004066	0. 56
45	NT2RP2004081	F-NT2RP2004081	0.94
	NT2RP2004098	F-NT2RP2004098	0. 39
	NT2RP2004124	F-NT2RP2004124	0.75
	NT2RP2004152	F-NT2RP2004152	0.54
50	NT2RP2004165	F-NT2RP2004165	0.68
	NT2RP2004187	F-NT2RP2004187	0.92
	NT2RP2004194	F-NT2RP2004194	0.90
	NT2RP2004196	F-NT2RP2004196	0.94
55	NT2RP2004226	F-NT2RP2004226	0.80
	NT2RP2004239	F-NT2RP2004239	0. 63

	NT2RP2004240	F-NT2RP2004240	0. 86
	NT2RP2004242	F-NT2RP2004242	0.94
	NT2RP2004245	F-NT2RP2004245	0.92
5	NT2RP2004316	F-NT2RP2004316_	0.33
	NT2RP2004364	F-NT2RP2004364	0.75
	NT2RP2004365	F-NT2RP2004365	0. 43
	NT2RP2004366	F-NT2RP2004366	0. 32
10	NT2RP2004373	F-NT2RP2004373	0. 60
	NT2RP2004389	F-NT2RP2004389	0. 94
	NT2RP2004392	F-NT2RP2004392	0. 93
	NT2RP2004399	F-NT2RP2004399	0.36
15	NT2RP2004463	F-NT2RP2004463	0. 43
	NT2RP2004476	F-NT2RP2004476	0. 80
	NT2RP2004538	F-NT2RP2004538	0. 94
	NT2RP2004551	F-NT2RP2004551	0.47
20	NT2RP2004568	F-NT2RP2004568	0.73
	NT2RP2004594	F-NT2RP2004594	0.64
	NT2RP2004600	F-NT2RP2004600	0. 53
	NT2RP2004602	F-NT2RP2004602	0.33
25	NT2RP2004614	F-NT2RP2004614	0.34
	NT2RP2004655	F-NT2RP2004655	0.77
	NT2RP2004664	F-NT2RP2004664	0. 94
	NT2RP2004689	F-NT2RP2004689	0.83
30	NT2RP2004710	F-NT2RP2004710	0.94
	NT2RP2004743	F-NT2RP2004743	0.32
	NT2RP2004768	F-NT2RP2004768	0.53
	NT2RP2004791	F-NT2RP2004791	0.64
35	NT2RP2004799	F-NT2RP2004799	0.42
33	NT2RP2004802	F-NT2RP2004802	0.88
	NT2RP2004816	F-NT2RP2004816	0.78
	NT2RP2004841	F-NT2RP2004841	0.61
40	NT2RP2004861	F-NT2RP2004861	0. 32
40	NT2RP2004897	F-NT2RP2004897	0.67
	NT2RP2004933	F-NT2RP2004933	0.94
	NT2RP2004936	F-NT2RP2004936	0. 53
	NT2RP2004959	F-NT2RP2004959	0. 78
45	NT2RP2004978	F-NT2RP2004978	0.86
	NT2RP2004985	F-NT2RP2004985	0. 42
	NT2RP2004999	F-NT2RP2004999	0.55
	NT2RP2005000	F-NT2RP2005000	0.40
50	NT2RP2005001	F-NT2RP2005001	0.94
	NT2RP2005003	F-NT2RP2005003	0.44
	NT2RP2005012 NT2RP2005018	F-NT2RP2005012 F-NT2RP2005018	0. 73 0. 43
	NT2RP2005018	F-NT2RP2005018	0. 43
55	NT2RP2005022	F-NT2RP2005037	0. 94
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	NT2RP2005038		0. 56
	NT2RP2005116		0. 94
	NT2RP2005126		0. 69
5	NT2RP2005139		0. 94
	NT2RP2005140	F-NT2RP2005140	0. 32
	NT2RP2005144	F-NT2RP2005144	0.39
	NT2RP2005147	F-NT2RP2005147	0. 32
10	NT2RP2005159	F-NT2RP2005159	0.52
	NT2RP2005162	F-NT2RP2005162	0.86
	NT2RP2005168	F-NT2RP2005168	0.53
	NT2RP2005204	F-NT2RP2005204	0.94
15	NT2RP2005227	F-NT2RP2005227	0. 53
	NT2RP2005239	F-NT2RP2005239	0. 91
	NT2RP2005270	F-NT2RP2005270	0.72
	NT2RP2005276	F-NT2RP2005276	0. 41
	NT2RP2005287	F-NT2RP2005287	0. 44
20	NT2RP2005288	F-NT2RP2005288	0. 82
	NT2RP2005315	F-NT2RP2005315	0. 79
	NT2RP2005358	F-NT2RP2005358	0. 94
	NT2RP2005393	F-NT2RP2005393	0. 41
25	NT2RP2005436	F-NT2RP2005436	0. 45
	NT2RP2005441	F-NT2RP2005441	0. 88
	NT2RP2005453	F-NT2RP2005453	0. 59
•	NT2RP2005457	F-NT2RP2005457	0, 56
30	NT2RP2005464	F-NT2RP2005464	0. 72
	NT2RP2005465	F-NT2RP2005465	0.40
	NT2RP2005472	F-NT2RP2005472	0.42
	NT2RP2005490	F-NT2RP2005490	0.52
35	NT2RP2005495	F-NT2RP2005495	0.94
	NT2RP2005496	F-NT2RP2005496	0.52
	NT2RP2005498	F-NT2RP2005498	0.74
	NT2RP2005509	F-NT2RP2005509	0. 55
40	NT2RP2005520	F-NT2RP2005520	0.54
	NT2RP2005525	F-NT2RP2005525	0.41
	NT2RP2005539	F-NT2RP2005539	0.94
	NT2RP2005540	F-NT2RP2005540	0.32
45	NT2RP2005549	F-NT2RP2005549	0.79
43	NT2RP2005555	F-NT2RP2005555	0.63
	NT2RP2005557	F-NT2RP2005557	0. 79
	NT2RP2005600	F-NT2RP2005600	0.76
	NT2RP2005605	F-NT2RP2005605	0.56
50 .	NT2RP2005620	F-NT2RP2005620	0.66
	NT2RP2005622	F-NT2RP2005622	0. 37
	NT2RP2005635	F-NT2RP2005635	0. 87
	NT2RP2005637	F-NT2RP2005637	0. 94
55	NT2RP2005640	F-NT2RP2005640	0. 87

	NT2RP2005654	F-NT2RP2005654	0.94
	NT2RP2005669	F-NT2RP2005669	0.94
	NT2RP2005675	F-NT2RP2005675	0.72
5	NT2RP2005683	F-NT2RP2005683	0.34
	NT2RP2005690	F-NT2RP2005690	0.54
	NT2RP2005712	F-NT2RP2005712	0.93
	NT2RP2005722	F-NT2RP2005722	0.50
10	NT2RP2005723	F-NT2RP2005723	0.45
	NT2RP2005732	F-NT2RP2005732	0.54
	NT2RP2005748	F-NT2RP2005748	0.76
	NT2RP2005752	F-NT2RP2005752	0.47
15	NT2RP2005763	F-NT2RP2005763	0.87
	NT2RP2005767	F-NT2RP2005767	0, 58
	NT2RP2005773	F-NT2RP2005773,	0.49
	NT2RP2005775	F-NT2RP2005775	0.34
20	NT2RP2005781	F-NT2RP2005781	0.81
20	NT2RP2005784	F-NT2RP2005784	0.94
	NT2RP2005804	F-NT2RP2005804	0.94
	NT2RP2005812	F-NT2RP2005812	0.90
-	NT2RP2005835	F-NT2RP2005835	0.94
25	NT2RP2005853	F-NT2RP2005853	0.77
	NT2RP2005859	F-NT2RP2005859	0.68
	NT2RP2005868	F-NT2RP2005868	0.76
	NT2RP2005886	F-NT2RP2005886	0.62
30	NT2RP2005890	F-NT2RP2005890	0.79
	NT2RP2005901	F-NT2RP2005901	0.94
	NT2RP2005933	F-NT2RP2005933	0.55
	NT2RP2006023	F-NT2RP2006023	0.47
35	NT2RP2006038	F-NT2RP2006038	0.73
	NT2RP2006043	F-NT2RP2006043	0. 50
	NT2RP2006052	F-NT2RP2006052	0.74
	NT2RP2006069	F-NT2RP2006069	0.85
40	NT2RP2006071	F-NT2RP2006071	0.94
	NT2RP2006100	F-NT2RP2006100	0.42
	NT2RP2006106	F-NT2RP2006106	0.94
	NT2RP2006141	F-NT2RP2006141	0.92
45	NT2RP2006186	F-NT2RP2006186	0. 94
	NT2RP2006196	F-NT2RP2006196	0. 32
	NT2RP2006200	F-NT2RP2006200	0. 70
	NT2RP2006219	F-NT2RP2006219	0.94
50	NT2RP2006237	F-NT2RP2006237	0. 59
	NT2RP2006238	F-NT2RP2006238	0. 47
	NT2RP2006275	F-NT2RP2006275	0.89
	NT2RP2006312	F-NT2RP2006312	0. 75
55	NT2RP2006333	F-NT2RP2006333	0. 73
	NT2RP2006334	F-NT2RP2006334	0. 72

	NT2RP2006365	F-NT2RP2006365	0.48
	NT2RP2006436	F-NT2RP2006436	0. 75
	NT2RP2006441	F-NT2RP2006441	0.61
5	NT2RP2006456	F-NT2RP2006456	0.60
	NT2RP2006464	F-NT2RP2006464	0. 51
	NT2RP2006472	F-NT2RP2006472	0. 76
	NT2RP2006565	F-NT2RP2006565	0. 34
10	NT2RP2006571	F-NT2RP2006571	0. 94
	NT2RP2006573	F-NT2RP2006573	0. 35
	NT2RP3000002	F-NT2RP3000002	0.40
	NT2RP3000031	F-NT2RP3000031	0.49
15	NT2RP3000046	F-NT2RP3000046	0.90
	NT2RP3000047	F-NT2RP3000047	0. 69
	NT2RP3000050	F-NT2RP3000050	0. 78
	NT2RP3000068	F-NT2RP3000068	0.94
20	NT2RP3000072	F-NT2RP3000072	0.34
	NT2RP3000080	F-NT2RP3000080	0. 32
	NT2RP3000085	F-NT2RP3000085	0.92
	NT2RP3000092	F-NT2RP3000092	0.40
25	NT2RP3000134	F-NT2RP3000134	0.42
	NT2RP3000142	F-NT2RP3000142	0.58
	NT2RP3000149	F-NT2RP3000149	0.83
•	NT2RP3000197	F-NT2RP3000197	0.59
00	NT2RP3000207	F-NT2RP3000207	0. 48
30	NT2RP3000220	F-NT2RP3000220	0. 75
	NT2RP3000251	F-NT2RP3000251	0.82
	NT2RP3000252	F-NT2RP3000252	0.88
	NT2RP3000255	F-NT2RP3000255	0. 94
35 .	NT2RP3000267	F-NT2RP3000267	0. 92
	NT2RP3000299	F-NT2RP3000299	0. 94
	NT2RP3000312	F-NT2RP3000312	0. 57
	NT2RP3000320	F-NT2RP3000320	0. 94
40	NT2RP3000324	F-NT2RP3000324	0.32
	NT2RP3000333	F-NT2RP3000333	0. 94
	NT2RP3000348	F-NT2RP3000348	0. 32
	NT2RP3000350	F-NT2RP3000350	0. 67
45	NT2RP3000359	F-NT2RP3000359	0.94
	NT2RP3000361	F-NT2RP3000361	0. 94
	NT2RP3000366	F-NT2RP3000366	0.94
	NT2RP3000393	F-NT2RP3000393	0.53
50	NT2RP3000397	F-NT2RP3000397	0.89
	NT2RP3000403	F-NT2RP3000403	0.87
	NT2RP3000439	F-NT2RP3000439	0.77
	NT2RP3000456	F-NT2RP3000456 F-NT2RP3000484	0. 88 0. 59
55	NT2RP3000484		0. 39
	NT2RP3000487	F-NT2RP3000487	u. 32

	NT2RP3000526	F-NT2RP3000526	0. 32
	NT2RP3000527	F-NT2RP3000527	0. 69
_	NT2RP3000531	F-NT2RP3000531	0.56
5	NT2RP3000562	F-NT2RP3000562	0.44
	NT2RP3000590	F-NT2RP3000590	0.88
	NT2RP3000592	F-NT2RP3000592	0.92
	NT2RP3000596	F-NT2RP3000596	0.50
10	NT2RP3000599	F-NT2RP3000599	0.69
	NT2RP3000603	F-NT2RP3000603	0.94
	NT2RP3000605	F-NT2RP3000605	0. 92
	NT2RP3000624	F-NT2RP3000624	0.71
15	NT2RP3000632	F-NT2RP3000632	0.43
	NT2RP3000644	F-NT2RP3000644	0.94
	NT2RP3000661	F-NT2RP3000661,	0.60
	NT2RP3000665	F-NT2RP3000665	0.46
20	NT2RP3000690	F-NT2RP3000690	0.79
·	NT2RP3000739	F-NT2RP3000739	0. 94
	NT2RP3000753	F-NT2RP3000753	0.42
	NT2RP3000759	F-NT2RP3000759	0.94
25	NT2RP3000825	F-NT2RP3000825	0.73
	NT2RP3000826	F-NT2RP3000826	0.94
	NT2RP3000836	F-NT2RP3000836	0.39
	NT2RP3000841	F-NT2RP3000841	0. 55
30	NT2RP3000845	F-NT2RP3000845	0. 38
30	NT2RP3000847	F-NT2RP3000847	0. 67
	NT2RP3000850	F-NT2RP3000850	0. 59
	NT2RP3000852	F-NT2RP3000852	0.39
	NT2RP3000859	F-NT2RP3000859	0. 83
35	NT2RP3000868	F-NT2RP3000868	0. 49
	NT2RP3000869	F-NT2RP3000869	0. 94
	NT2RP3000901	F-NT2RP3000901	0. 34
	NT2RP3000917	F-NT2RP3000917	0. 73
40	NT2RP3000919	F-NT2RP3000919	0. 94
	NT2RP3000968	F-NT2RP3000968	0. 94
	NT2RP3000980	F-NT2RP3000980	0. 61
	NT2RP3000994	F-NT2RP3000994	0. 45
45	NT2RP3001004	F-NT2RP3001004	0. 86
	NT2RP3001055	F-NT2RP3001055	0. 38
	NT2RP3001057	F-NT2RP3001057	0. 84
	NT2RP3001081	F-NT2RP3001081	0. 57
50	NT2RP3001084	F-NT2RP3001084	0.94
	NT2RP3001096	F-NT2RP3001096	0. 94
	NT2RP3001107	F-NT2RP3001107	0. 88
	NT2RP3001109	F-NT2RP3001109	0. 67
55	NT2RP3001111	F-NT2RP3001111	0.87
	NT2RP3001113	F-NT2RP3001113	0. 40

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	NT2RP3001116	F-NT2RP3001116	0. 73
	NT2RP3001119	F-NT2RP3001119	0. 62
•	NT2RP3001120	F-NT2RP3001120	0.59
5	NT2RP3001126	F-NT2RP3001126	0.94
	NT2RP3001133	F-NT2RP3001133	0.89
	NT2RP3001140	F-NT2RP3001140	0. <b>59</b>
	NT2RP3001147	F-NT2RP3001147	0. 94
10	NT2RP3001150	F-NT2RP3001150	0.92
	NT2RP3001155	F-NT2RP3001155	0.94
	NT2RP3001176	F-NT2RP3001176	0.82
	NT2RP3001214	F-NT2RP3001214	0.41
15	NT2RP3001216	F-NT2RP3001216	0.32
	NT2RP3001221	F-NT2RP3001221	0.46
	NT2RP3001236	F-NT2RP3001236,	0.79
	NT2RP3001239	F-NT2RP3001239	0.94
20	NT2RP30G1253	F-NT2RP3001253	0.59
	NT2RP3001260	F-NT2RP3001260	0.94
	NT2RP3001268	F-NT2RP3001268	0.75
	NT2RP3001272	F-NT2RP3001272	0.94
25	NT2RP3001274	F-NT2RP3001274	0.94
20	NT2RP3001297	F-NT2RP3001297	0.94
	NT2RP3001307	F-NT2RP3001307	0.87
	NT2RP3001325	F-NT2RP3001325	0.71
	NT2RP3001338	F-NT2RP3001338	0.90
30	NT2RP3001355	F-NT2RP3001355	0.40
	NT2RP3001356	F-NT2RP3001356	0.88
	NT2RP3001384	F-NT2RP3001384	0.91
	NT2RP3001392	F-NT2RP3001392	0.42
35	NT2RP3001396	F-NT2RP3001396	0.56
	NT2RP3001398	F-NT2RP3001398	0.94
	NT2RP3001399	F-NT2RP3001399	0. 94
	NT2RP3001407	F-NT2RP3001407	0. 54
40	NT2RP30C1420	F-NT2RP3001420	0.74
	NT2RP3001426	F-NT2RP3001426	0. 92
	NT2RP3001427	F-NT2RP3001427	0.40
	NT2RP3001428	F-NT2RP3001428	0.94
45	NT2RP3001447	F-NT2RP3001447	0.57
	NT2RP3001449	F-NT2RP3001449	0.62
	NT2RP3001453	F-NT2RP3001453	0.94
	NT2RP3001457	F-NT2RP3001457	0.89
50	NT2RP3001472	F-NT2RP3001472	0.82
<del></del>	NT2RP3001490	F-NT2RP3001490	0. 53
	NT2RP3001495	F-NT2RP3001495	0. 94
	NT2RP3001497	F-NT2RP3001497	0.40
55	NT2RP3001529	F-NT2RP3001529	0.94
55	NT2RP3001538	F-NT2RP3001538	0. 93

	NT-2RP3001554	F-NT2RP3001554	0.91
	NT2RP3001587	F-NT2RP3001587	0.94
	NT2RP3001608	F-NT2RP3001608	0.94
5	NT2RP3001621	F-NT2RP3001621	0.78
	NT2RP3001629	F-NT2RP3001629	0.42
	NT2RP3001642	F-NT2RP3001642	0.94
	NT2RP3001646	F-NT2RP3001646	0.94
10	NT2RP3001671	F-NT2RP3001671	0.77
	NT2RP3001672	F-NT2RP3001672	0.88
	NT2RP3001676	F-NT2RP3001676	0.88
	NT2RP3001678	F-NT2RP3001678	0.44
15	NT2RP3001679	F-NT2RP3001679	0.44
	NT2RP3001688	F-NT2RP3001688	0.67
	NT2RP3001690	F-NT2RP3001690	0.58
	NT2RP3001698	F-NT2RP3001698	0.86
20	NT2RP3001708	F-NT2RP3001708	0.38
	NT2RP3001712	F-NT2RP3001712	0.75
	NT2RP3001716	F-NT2RP3001716	0.77
	NT2RP3001724	F-NT2RP3001724	0.65
25	NT2RP3001727	F-NT2RP3001727	0.84
	NT2RP3001730	F-NT2RP3001730	0.77
	NT2RP3001739	F-NT2RP3001739	0. 78
	NT2RP3001753	F-NT2RP3001753	0. 79
30	NT2RP3001764	F-NT2RP3001764	0.33
	NT2RP3001777	F-NT2RP3001777	0.87
•	NT2RP3001792	F-NT2RP3001792	0.86
	NT2RP3001799	F-NT2RP3001799	0. 79
	NT2RP3001819	F-NT2RP3001819	0.31
35	NT2RP3001844	F-NT2RP3001844	0.34
	NT2RP3001854	F-NT2RP3001854	0.94
	NT2RP3001855	F-NT2RP3001855	0.88
	NT2RP3001857	F-NT2RP3001857	0.41
40	NT2RP3001896	F-NT2RP3001896	0.94
	NT2RP3001915		0.42
	NT2RP3001929	F-NT2RP3001929	0.49
	NT2RP3001931	F-NT2RP3001931	0.94
45	NT2RP3001938	F-NT2RP3001938	0.32
	NT2RP3001943	F-NT2RP3001943	0.70
	NT2RP3001944 NT2RP3001969		0. 94 0. 54
	NT2RP3002007		0.54
50	NT2RP3002007		0.94
	NT2RP3002014 NT2RP3002033		0.94
	NT2RP3002033		0. 94
	NT2RP3002045		0. 94
55	NT2RP3002062		0. 62
	412N 3002002	1 MIZM 0002002	0. 02

	NT2RP3002063	F-NT2RP3002063	0.34
	NT2RP3002081	F-NT2RP3002081	0.57
	NT2RP3002097	F-NT2RP3002097	0. 72
5	NT2RP3002102	F-NT2RP3002102	0. 90
	NT2RP3002102	F-NT2RP3002102	0.74
		F-NT2RP3002142	0. 74
	NT2RP3002142		
	NT2RP3002146	F-NT2RP3002146	0.91
10	NT2RP3002147	F-NT2RP3002147	0.37
	NT2RP3002151	F-NT2RP3002151	0.90
	NT2RP3002163	F-NT2RP3002163	0.76
	NT2RP3002165	F-NT2RP3002165	0.89
15	NT2RP3002181	F-NT2RP3002181	0.78
	NT2RP3002244	F-NT2RP3002244	0.81
	NT2RP3002248	F-NT2RP3002248,	
	NT2RP3002255	F-NT2RP3002255	0.53
20	NT2RP3002273	F-NT2RP3002273	0.90
	NT2RP3002276	F-NT2RP3002276	0.39
	NT2RP3002303	F-NT2RP3002303	0.40
	NT2RP3002304	F-NT2RP3002304	0.44
25	NT2RP3002330	F-NT2RP3002330	0.94
25	NT2RP3002343	F-NT2RP3002343	0.94
	NT2RP3002351	F-NT2RP3002351	0.94
	NT2RP3002377	F-NT2RP3002377	0.52
	NT2RP3002399	F-NT2RP3002399	0. 91
30	NT2RP3002402	F-NT2RP3002402	0. 33
	NT2RP3002455	F-NT2RP3002455	0.42
	NT2RP3002484	F-NT2RP3002484	0.52
	NT2RP3002501	F-NT2RP3002501	0.57
35	NT2RP3002512	F-NT2RP3002512	0.63
	NT2RP3002529	F-NT2RP3002529	0. 74
	NT2RP3002545	F-NT2RP3002545	0.94
	NT2RP3002549	F-NT2RP3002549	0.94
40	NT2RP3002566	F-NT2RP3002566	0.74
	NT2RP3002587	F-NT2RP3002587	0.94
	NT2RP3002590	F-NT2RP3002590	0.37
	NT2RP3002602	F-NT2RP3002602	0.39
45	NT2RP3002603	F-NT2RP3002603	0.46
•	NT2RP3002628	F-NT2RP3002628	0.31
	NT2RP3002631	F-NT2RP3002631	0.79
	NT2RP3002650	F-NT2RP3002650	0.94
	NT2RP3002659	F-NT2RP3002659	0.93
50	NT2RP3002660	F-NT2RP3002660	0.46
	NT2RP3002663	F-NT2RP3002663	0.79
	NT2RP3002671	F-NT2RP3002671	0.34
	NT2RP3002682	F-NT2RP3002682	0.52
55	NT2RP3002688	F-NT2RP3002688	0.82

	NT2RP3002701	F-NT2RP3002701	0. 62
	NT2RP3002763	F-NT2RP3002763	0. 83
	NT2RP3002785	F-NT2RP3002785	0. 33
5	NT2RP3002818	F-NT2RP3002818	
	NT2RP3002861	F-NT2RP3002861	0.31
	NT2RP3002869	F-NT2RP3002869	0. 47
	NT2RP3002876	F-NT2RP3002876	0. 58
10	NT2RP3002877	F-NT2RP3002877	0. 33
	NT2RP3002977	F-NT2RP3002909	0. 52
		F-NT2RP3002909	0. 52
	NT2RP3002911		
	NT2RP3002948	F-NT2RP3002948	0.60
15	NT2RP3002953	F-NT2RP3002953	0. 37
	NT2RP3002969	F-NT2RP3002969	0. 47
	NT2RP3002972	F-NT2RP3002972,	
	NT2RP3002985	F-NT2RP3002985	0.55
20	NT2RP3002988	F-NT2RP3002988	0.82
	NT2RP3003008	F-NT2RP3003008	0. 94
	NT2RP3003061	F-NT2RP3003061	0. 93
	NT2RP3003068	F-NT2RP3003068	0. 45
25	NT2RP3003071	F-NT2RP3003071	0.76
	NT2RP3003078	F-NT2RP3003078	0.40
	NT2RP3003101	F-NT2RP3003101	0.41
	NT2RP3003139	F-NT2RP3003139	0.43
30	NT2RP3003145	F-NT2RP3003145	0. 94
	NT2RP3003150	F-NT2RP3003150	0. 94
	NT2RP3003157	F-NT2RP3003157	0. 81
	NT2RP3003185	F-NT2RP3003185	0. 87
	NT2RP3003193	F-NT2RP3003193	0. 94
35	NT2RP3003197	F-NT2RP3003197	0. 74
	NT2RP3003203	F-NT2RP3003203	0.94
	NT2RP3003204	F-NT2RP3003204	0. 74
	NT2RP3003210	F-NT2RP3003210	0.45
40	NT2RP3003212	F-NT2RP3003212	0.51
	NT2RP3003230	F-NT2RP3003230	0. 94
	NT2RP3003242	F-NT2RP3003242	0.92
	NT2RP3003251	F-NT2RP3003251	0. 93
45	NT2RP3003278	F-NT2RP3003278	0.83
	NT2RP3003282	F-NT2RP3003282	0. 67
	NT2RP3003290	F-NT2RP3003290	0.62
	NT2RP3003301	F-NT2RP3003301	0. 49
50	NT2RP3003302	F-NT2RP3003302	0.32
	NT2RP3003311	F-NT2RP3003311	0.34
	NT2RP3003313	F-NT2RP3003313	0.88
	NT2RP3003327	F-NT2RP3003327	0.88
55	NT2RP3003330	F-NT2RP3003330	0. 94
33	NT2RP3003344	F-NT2RP3003344	0.67

	NT2RP3003353	F-NT2RP3003353	0. 65
	NT2RP3003377	F-NT2RP3003377	0.87
	NT2RP3003385	F-NT2RP3003385	0.94
5	NT2RP3003409	F-NT2RP3003409	0.41
	NT2RP3003411	F-NT2RP3003411	0. 47
	NT2RP3003427	F-NT2RP3003427	0. 33
	NT2RP3003427	F-NT2RP3003490	0.56
		F-NT2RP3003491	0.37
10	NT2RP3003491		0. 37
	NT2RP3003543	F-NT2RP3003543	
	NT2RP3003552	F-NT2RP3003552	0.64
	NT2RP3003555	F-NT2RP3003555	0.94
15	NT2RP3003564	F-NT2RP3003564	0.86
	NT2RP3003589	F-NT2RP3003589	0.88
	NT2RP3003621	F-NT2RP3003621,	
	NT2RP3003656	F-NT2RP3003656	0. 77
20	NT2RP3003659	F-NT2RP3003659	0. 32
	NT2RP3003672	F-NT2RP3003672	0.49
	NT2RP3003680	F-NT2RP3003680	0.94
	NT2RP3003686	F-NT2RP3003686	0. 78
25	NT2RP3003701	F-NT2RP3003701	0.36
25	NT2RP3003716	F-NT2RP3003716	0.34
	NT2RP3003726	F-NT2RP3003726	0. 72
	NT2RP3003795	F-NT2RP3003795	0. 34
	NT2RP3003805	F-NT2RP3003805	0.74
30	NT2RP3003809	F-NT2RP3003809	0. 73
	NT2RP3003825	F-NT2RP3003825	0.94
	NT2RP3003831	F-NT2RP3003831	0. 52
	NT2RP3003833	F-NT2RP3003833	0.31
35	NT2RP3003846	F-NT2RP3003846	0.54
	NT2RP3003870	F-NT2RP3003870	0.91
	NT2RP3003876	F-NT2RP3003876	0.42
	NT2RP3003914	F-NT2RP3003914	0.94
40	NT2RP3003918	F-NT2RP3003918	0.90
•	NT2RP3003932	F-NT2RP3003932	0.64
	NT2RP3003992	F-NT2RP3003992	0. 94
	NT2RP3004013	F-NT2RP3004013	0. 60
	NT2RP3004041	F-NT2RP3004041	0, 52
45	NT2RP3004051	F-NT2RP3004051	0. 75
	NT2RP3004078	F-NT2RP3004078	0. 73
	NT2RP3004093	F-NT2RP3004093	0. 64
	NT2RP3004125	F-NT2RP3004125	0. 94
50	NT2RP3004155	F-NT2RP3004155	0. 89
	NT2RP3004189	F-NT2RP3004189	0. 46
	NT2RP3004189	F-NT2RP3004206	0. 59
	NT2RP3004206	F-NT2RP3004206	0. 39
55		F-NT2RP3004207	0. 54
	NT2RP3004209	F-N12NF3004209	V. 54

	NT2RP3004242	F-NT2RP3004242	0. 61
	NT2RP3004246	F-NT2RP3004246	0. 91
	NT2RP3004253	F-NT2RP3004253	0.41
5	NT2RP3004258	F-NT2RP3004258	
	NT2RP3004262	F-NT2RP3004262	0.56
	NT2RP3004282	F-NT2RP3004282	0, 90
	NT2RP3004332	F-NT2RP3004332	0. 53
10	NT2RP3004341	F-NT2RP3004341	0.40
	NT2RP3004348	F-NT2RP3004348	0.94
	NT2RP3004378	F-NT2RP3004378	0.40
	NT2RP3004424	F-NT2RP3004424	0. 75
15	NT2RP3004428	F-NT2RP3004428	0.85
	NT2RP3004451	F-NT2RP3004451	0.85
	NT2RP3004454	F-NT2RP3004454,	
	NT2RP3004466	F-NT2RP3004466	0.80
00	NT2RP3004472	F-NT2RP3004472	0.46
20	NT2RP3004480	F-NT2RP3004480	0. 77
	NT2RP3004490	F-NT2RP3004490	0. 65
	NT2RP3004498	F-NT2RP3004498	0. 91
	NT2RP3004504	F-NT2RP3004504	0, 88.
25	NT2RP3004507	F-NT2RP3004507	0.86
	NT2RP3004534	F-NT2RP3004534	0. 32
	NT2RP3004539	F-NT2RP3004539	0.76
	NT2RP3004544	F-NT2RP3004544	0. 94
30	NT2RP3004566	F-NT2RP3004566	0.94
	NT2RP3004569	F-NT2RP3004569	0.69
	NT2RP3004572	F-NT2RP3004572	0.65
	NT2RP3004578	F-NT2RP3004578	0.52
35	NT2RP3004594	F-NT2RP3004594	0.39
	NT2RP3004617	F-NT2RP3004617	0.94
	NT2RP3004618	F-NT2RP3004618	0.41
	NT2RP3004669	F-NT2RP3004669	0. 82
40	NT2RP3004670	F-NT2RP3004670	0. 32
	NT2RP4000008	F-NT2RP4000008	0.94
	NT2RP4000051	F-NT2RP4000051	0.94
	NT2RP4000109	F-NT2RP4000109	0.84
45	NT2RP4000111	F-NT2RP4000111	0.49
	NT2RP4000129	F-NT2RP4000129	0. 94
	NT2RP4000147	F-NT2RP4000147	0. 92
	NT2RP4000150	F-NT2RP4000150	0. 41
50	NT2RP4000151	F-NT2RP4000151	0. 33
	NT2RP4000159	F-NT2RP4000159	0. 75
	NT2RP4000185	F-NT2RP4000185	0. 94
	NT2RP4000210	F-NT2RP4000210	0.83
55	NT2RP4000212		0. 94
55	NT2RP4000243	F-NT2RP4000243	0. 94

	NT2RP4000246-	F-NT2RP4000246	0. 59
	NT2RP4000259	F-NT2RP4000259	0. 94
	NT2RP4000263	F-NT2RP4000263	0. 62
5	NT2RP4000290	F-NT2RP4000290	0. 92
	NT2RP4000312	F-NT2RP4000312	0. 39
	NT2RP4000323	F-NT2RP4000323	0.73
	NT2RP4000355	F-NT2RP4000355	0. 94
10	NT2RP4000360	F-NT2RP4000360	0. 55
	NT2RP4000367	F-NT2RP4000367	0. 60
	NT2RP4000370	F-NT2RP4000370	0.93
	NT2RP4000376	F-NT2RP4000376	0.88
15	NT2RP4000381	F-NT2RP4000381	0.39
	NT2RP4000398	F-NT2RP4000398	0.64
	NT2RP4000415	F-NT2RP4000415,	0.94
	NT2RP4000417	F-NT2RP4000417	0. 33
20	NT2RP4000448	F-NT2RP4000448	0.33
	NT2RP4000449	F-NT2RP4000449	033
	NT2RP4000455	F-NT2RP4000455	0.37
	NT2RP4000457	F-NT2RP4000457	0.36
ar.	NT2RP4000480	F-NT2RP4000480	0.88
25	NT2RP4000481	F-NT2RP4000481	0.51
	NT2RP4000498	F-NT2RP4000498	0.64
	NT2RP4000500	F-NT2RP4000500	0.67
	NT2RP4000518	F-NT2RP4000518	0.34
30	NT2RP4000524	F-NT2RP4000524	0. 94
,	NT2RP4000528	F-NT2RP4000528	0. 65
	NT2RP4000541	F-NT2RP4000541	0. 44
	NT2RP4000556	F-NT2RP4000556	0.69
35	NT2RP4000560	F-NT2RP4000560	0. 67
	NT2RP4000588	F-NT2RP4000588	0.94
	NT2RP4000614	F-NT2RP4000614	0. 94
	NT2RP4000638	F-NT2RP4000638	0. 74
40	NT2RP4000648	F-NT2RP4000648	0. 40
	NT2RP4000657	F-NT2RP4000657	0. 94
	NT2RP4000704		0. 69
	NT2RP4000713	F-NT2RP4000713	0. 93
45	NT2RP4000724	F-NT2RP4000724	
	NT2RP4000728		0. 34
	NT2RP4000737		0. 48
	NT2RP4000739		0.40
50	NT2RP4000781	F-NT2RP4000781	0.94
•	NT2RP4000787		0.94
	NT2RP4000817		0.60
	NT2RP4000833		0.35
55	NT2RP4000837		0.80
	NT2RP4000839	F-NT2RP4000839	0. 33

	NT2RP4000855	F-NT2RP4000855	0. 74
	NT2RP4000865	F-NT2RP4000865	0. 77
	NT2RP4000878	F-NT2RP4000878	0. 79
5	NT2RP4000879	F-NT2RP4000879	0. 41
	NT2RP4000907	F-NT2RP4000907	0. 44
	NT2RP4000918	F-NT2RP4000918	0. 88
	NT2RP4000925	F-NT2RP4000925	0. 89
10	NT2RP4000927	F-NT2RP4000927	0. 81
	NT2RP4000928	F-NT2RP4000928	0.39
	NT2RP4000929	F-NT2RP4000929	0.34
	NT2RP4000955	F-NT2RP4000955	0. 77
15	NT2RP4000973	F-NT2RP4000973	0.36
	NT2RP4000975	F-NT2RP4000975	0. 87
	NT2RP4000979	F-NT2RP4000979,	0.50
	NT2RP4000984	F-NT2RP4000984	0.66
20	NT2RP4000989	F-NT2RP4000989	0. 94
20	NT2RP4000997	F-NT2RP4000997	0.68
	NT2RP4001004	F-NT2RP4001004	0.40
	NT2RP4001006	F-NT2RP4001006	0.91
	NT2RP4001010	F-NT2RP4001010	0.64
25	NT2RP4001029	F-NT2RP4001029	0.77
	NT2RP4001041	F-NT2RP4001041	0.79
	NT2RP4001057	F-NT2RP4001057	0. 65
	NT2RP4001064	F-NT2RP4001064	0.94
30	NT2RP4001078	F-NT2RP4001078	0. 39
	NT2RP4001079	F-NT2RP4001079	0. 94
	NT2RP40G1080	F-NT2RP4001080	0. 58
	NT2RP4001086	F-NT2RP4001086	0. 78
35	NT2RP4001095	F-NT2RP4001095	0. 80
	NT2RP4001100	F-NT2RP4001100	0. 79
	NT2RP4001117	F-NT2RP4001117	0. 60
	NT2RP4001122	F-NT2RP4001122	0. 82
40	NT2RP4001126	F-NT2RP4001126	0. 55
	NT2RP4001138		0. 94
	NT2RP4001143		0, 41 0, 37
	NT2RP4001148		
45	N12RP4001149		0.50
	NT2RP4001150		0. 94 0. 78
	NT2RP4001159		0. 78
	NT2RP4001174		0. 80
50	NT2RP4001206 NT2RP4001207		0. 38
	NT2RP4001207		
	NT2RP4001210		
	NT2RP4001213		
55	NT2RP4001219	•	
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	NT2RP4001235	F-NT2RP4001235	0.88
	NT2RP4001256	F-NT2RP4001256	0.60
	NT2RP4001260	F-NT2RP4001260	0. 94
5	NT2RP4001274	F-NT2RP4001274	0.82
	NT2RP4001276	F-NT2RP4001276	0. 68
	NT2RP4001313	F-NT2RP4001313	0.77
	NT2RP4001315	F-NT2RP4001315	0.87
10	NT2RP4001336	F-NT2RP4001336	0.48
	NT2RP4001339	F-NT2RP4001339	0.41
·	NT2RP4001343	F-NT2RP4001343	0. 63
	NT2RP4001345	F-NT2RP4001345	0. 89
15	NT2RP4001351	F-NT2RP4001351	0. 87
	NT2RP4001353	F-NT2RP4001353	0. 72
	NT2RP4001372	F-NT2RP4001372,	• • • •
	NT2RP4001373	F-NT2RP4001373	0.60
22	NT2RP4001375	F-NT2RP4001375	0. 43
20	NT2RP4001379	F-NT2RP4001379	0.77
	NT2RP4001389	F-NT2RP4001389	0. 41
	NT2RP4001414	F-NT2RP4001414	0. 90
	NT2RP4001433	F-NT2RP4001433	0.39
25	NT2RP4001442	F-NT2RP4001442	0.59
	NT2RP4001447	F-NT2RP4001447	0. 33
	NT2RP4001474	F-NT2RP4001474	0.89
	NT2RP4001483	F-NT2RP4001483	0. 94
30	NT2RP4001498	F-NT2RP4001498	0. 59
	NT2RP4001502	F-NT2RP4001502	0.61
	NT2RP4001507	F-NT2RP4001507	0. 91
	NT2RP4001524	F-NT2RP4001524	0.61
35	NT2RP4001529	F-NT2RP4001529	0.75
	NT2RP4001547	F-NT2RP4001547	0.77
	NT2RP4001551	F-NT2RP4001551	0.79
	NT2RP4001555	F-NT2RP4001555	0. 79
40	NT2RP4001567	F-NT2RP4001567	0.39
	NT2RP4001568	F-NT2RP4001568	0, 45
	NT2RP4001571	F-NT2RP4001571	0. 76
	NT2RP4001574	F-NT2RP4001574	0. 91
45	NT2RP4001575	F-NT2RP4001575	0.57
40	NT2RP4001592	F-NT2RP4001592	0.71
	NT2RP4001610	F-NT2RP4001610	0. 94
	NT2RP4001614	F-NT2RP4001614	0.73
	NT2RP4001634	F-NT2RP4001634	0.40
50	NT2RP4001638	F-NT2RP4001638	0. 37
	NT2RP4001644	F-NT2RP4001644	0.94
	NT2RP4001656	F-NT2RP4001656	0. 92
	NT2RP4001677	F-NT2RP4001677	0. 47
55	NT2RP4001679	F-NT2RP4001679	0. 54
			•

	NT2RP4001696	F-NT2RP4001696	0. 62
	NT2RP4001725	F-NT2RP4001725	0. 94
	NT2RP4001730	F-NT2RP4001730	0. 94
5	NT2RP4001739	F-NT2RP4001739	0.82
	NT2RP4001753	F-NT2RP4001753	0. 66
	NT2RP4001760	F-NT2RP4001760	0. 39
	NT2RP4001790	F-NT2RP4001790	0.64
10	NT2RP4001803	F-NT2RP4001803	0. 87
	NT2RP4001822	F-NT2RP4001822	0.84
	NT2RP4001823	F-NT2RP4001823	0.56
	NT2RP4001828	F-NT2RP4001828	0.45
15	NT2RP4001838	F-NT2RP4001838	0.87
	NT2RP4001841	F-NT2RP4001841	0.58
	NT2RP4001849	F-NT2RP4001849,	0.88
	NT2RP4001861	F-NT2RP4001861	0.80
20	NT2RP4001893	F-NT2RP4001893	0.94
	NT2RP4001896	F-NT2RP4001896	0.93
	NT2RP4001901	F-NT2RP4001901	0.89
	NT2RP4001927	F-NT2RP4001927	0.77
25	NT2RP4001938	F-NT2RP4001938	0.94
23	NT2RP4001946	F-NT2RP4001946	0.74
	NT2RP4001950	F-NT2RP4001950	0.45
	NT2RP4001953	F-NT2RP4001953	0.49
20	NT2RP4001966	F-NT2RP4001966	0.86
30	NT2RP4001975	F-NT2RP4001975	0.94
	NT2RP4002018	F-NT2RP4002018	0.67
	NT2RP4002047	F-NT2RP4002047	0.67
	NT2RP4002052	F-NT2RP4002052	0.42
35	NT2RP4002058	F-NT2RP4002058	0. 78
	NT2RP4002071	F-NT2RP4002071	0.94
	NT2RP4002078	F-NT2RP4002078	0.38
	NT2RP4002081	F-NT2RP4002081	0.55
40	NT2RP4002083	F-NT2RP4002083	0.86
	NT2RP4002408	F-NT2RP4002408	0.64
	NT2RP4002791	F-NT2RP4002791	0.84
	NT2RP4002888	F-NT2RP4002888	0.31
45	NT2RP5003459		
•	NT2RP5003461	F-NT2RP5003461	0.94
	NT2RP5003477	F-NT2RP5003477	0.44
	NT2RP5003492	F-NT2RP5003492	0.57
50	NT2RP5003500	F-NT2RP5003500	0.94
	NT2RP5003506	F-NT2RP5003506 F-NT2RP5003512	0.94
	NT2RP5003512 NT2RP5003522		0. 42 0. 94
	NT2RP5003522 NT2RP5003524	F-NT2RP5003522 F-NT2RP5003524	0. 58
55	NT2RP5003524	F-NT2RP5003534	0. 32
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	OVARC1000001	F-0VARC1000001	0. 74
		F-0VARC1000004	0. 90
	OVARC1000006	F-0VARC1000006	0.94
5	OVARC1000013	F-0VARC1000013	0.48
	OVARC1000014	F-0VARC1000014	0.69
	OVARC1000035	F-0VARC1000035	0.59
	OVARC1000060	F-0VARC1000060	0.76
10	OVARC1000087	F-0VARC1000087	0.31
	OVARC1000091	F-0VARC1000091	0.88
	OVARC1000106	F-0VARC1000106	0.88
	OVARC1000109	F-0VARC1000109	0.94
15	OVARC1000113	F-0VARC1000113	0.40
	OVARC1000114	F-0VARC1000114	0.53
	OVARC1000139	F-0VARC1000139,	0.88
	OVARC1000148	F-0VARC1000148	0.94
20	0VARC1000151	F-0VARC1000151	0.34
20	OVARC1000168	F-0VARC1000168	0.56
	OVARC1000209	F-0VARC1000209	0.94
	OVARC1000212	F-0VARC1000212	0.87
	OVARC1000241	F-0VARC1000241	0.59
25	OVARC1000288	F-0VARC1000288	0. 87
	OVARC1000304	F-0VARC1000304	0.94
	OVARC1000309	F-0VARC1000309	0.76
	OVARC1000321	F-0VARC1000321	0.90
30	OVARC1000326	F-0VARC1000326	0.46
	OVARC1000335	F-0VARC1000335	0.55
	OVARC1000347	F-0VARC1000347	0.43
	OVARC1000384	F-0VARC1000384	0.90
35	OVARC1000408	F-0VARC1000408	0.42
	OVARC1000411	F-0VARC1000411	0.65
	OVARC1000420	F-0VARC1000420	0.84
	OVARC1000437	F-0VARC1000437	0.94
40	OVARC1000440	F-0VARC1000440	0.94
	OVARC1000442	F-0VARC1000442	0.94
	OVARC1000443	F-0VARC1000443	0.74
	0VARC1000461	F-0VARC1000461	0.89
45	OVARC1000465	F-0VARC1000465	0.75
40	OVARC1000466	F-0VARC1000466	0.94
	OVARC1000473	F-0VARC1000473	0.31
	OVARC1000479	F-0VARC1000479	0.42
	OVARC1000520	F-0VARC1000520	0. 94
50	OVARC1000556	F-0VARC1000556	0.94
	OVARC1000557	F-0VARC1000557	0.72
	OVARC1000564	F-0VARC1000564	0.94
	0VARC1000576	F-0VARC1000576	0.40
55	OVARC1000588	F-0VARC1000588	0. 45

	OVARC1000605	F-0VARC1000605	0.58
	OVARC1000622	F-0VARC1000622	0.80
	OVARC1000640	F-0VARC1000640	0.35
5	OVARC1000649	F-0VARC1000649	0.94
	OVARC1000661	F-0VARC1000661	0.87
	0VARC1000681	F-0VARC1000681	0.84
	OVARC1000682	F-0VARC1000682	0.38
10	OVARC1000689	F-0VARC1000689	0.36
	OVARC1000703	F-0VARC1000703	0.92
	OVARC1000722	F-0VARC1000722	0.94
	OVARC1000730	F-0VARC1000730	0.90
15	OVARC1000746	F-0VARC1000746	0.42
	OVARC1000771	F-0VARC1000771	0.94
·	OVARC1000781	F-0VARC1000781,	0.80
	OVARC1000787	F-0VARC1000787	0.41
20	OVARC1000800	F-0VARC1000800	0.85
20	OVARC1000834	F-0VARC1000834	0.62
	OVARC1000846	F-0VARC1000846	0.49
	OVARC1000850	F-0VARC1000850	0.94
	OVARC1000862	F-0VARC1000862	0.54
25	OVARC1000876	F-0VARC1000876	0.74
	OVARC1000883	F-0VARC1000883	0.55
	OVARC1000885	F-0VARC1000885	0.94
	OVARC1000886	F-0VARC1000886	0.61
30	OVARC1000890	F-0VARC1000890	0.45
	OVARC1000912	F-0VARC1000912	0.63
	0VARC1000915	F-0VARC1000915	0.31
•	OVARC1000924	F-0VARC1000924	0.82
35	0VARC1000945	F-0VARC1000945	0.74
	0VARC1000959	F-0VARC1000959	0.33
	OVARC1000964	F-0VARC1000964	0.60
	0VARC1000984	F-0VARC1000984	0.59
40	0VARC1000996	F-0VARC1000996	0, 83
	0VARC1000999	F-0VARC1000999	0.94
	0VARC1001004	F-0VARC1001004	0.69
	0VARC1001010	F-0VARC1001010	0.51
45	0VARC1001032	F-0VARC1001032	0.32
	0VARC1001034	F-0VARC1001034	0. 78
	0VARC1001038	F-0VARC1001038	0. 93
	OVARC1001040	F-0VARC1001040	0.68
50	0VARC1001044	F-0VARC1001044	0.66
	0VARC1001055	F-0VARC1001055	0. 76
	OVARC1001065	F-0VARC1001065	0.31
	0VARC1001068	F-0VARC1001068	0.63
55	0VARC1001074	F-0VARC1001074	0.82
	0VARC1001092	F-0VARC1001092	0. 50

	OVARC1001107	F-0VARC1001107	0.65
	OVARC1001113	F-0VARC1001113	0.86
	OVARC1001154	F-0VARC1001154	0.43
5	OVARC1001161	F-0VARC1001161	0, 75
	OVARC1001162	F-0VARC1001162	0.61
	OVARC1001167	F-0VARC1001167	0.31
	OVARC1001170	F-0VARC1001170	0.46
10	OVARC1001171	F-0VARC1001171	0.78
	OVARC1001173	F-0VARC1001173	0.71
	OVARC1001176	F-0VARC1001176	0.83
	OVARC1001180	F-0VARC1001180	0.53
15	OVARC1001188	F-0VARC1001188	0.42
	OVARC1001200	F-0VARC1001200	0.94
	OVARC1001232	F-0VARC1001232;	0.90
	OVARC1001243	F-0VARC1001243	0.59
20	OVARC1001244	F-0VARC1001244	0.79
	OVARC1001270	F-0VARC1001270	0.48
	OVARC1001271	F-0VARC1001271	0.67
	OVARC1001296	F-0VARC1001296	0.60
25	OVARC1001306	F-0VARC1001306	0.44
25	OVARC1001329	F-0VARC1001329	0. 91
	OVARC1001341	F-0VARC1001341	0.82
	OVARC1001344	F-0VARC1001344	0. 94
	OVARC1001360	F-0VARC1001360	0. 94
30	OVARC1001369	F-0VARC1001369	0.36
	OVARC1001372	F-0VARC1001372	0. 63
	OVARC1001376	F-0VARC1001376	0.81
	OVARC1001381	F-0VARC1001381	0. 78
35	0VARC1001391	F-0VARC1001391	0. 33
	OVARC1001399	F-0VARC1001399	0. 67
	OVARC1001417	F-0VARC1001417	0.94
	0VARC1001419	F-0VARC1001419	0. 34 0. 73
40	0VARC1001425	F-0VARC1001425 F-0VARC1001436	0. 73
	OVARC1001436 OVARC1001453	F-0VARC1001453	0. 45
	0VARC1001476	F-0VARC1001435	0. 45
	0VARC1001470		0. 35
45	0VARC1001496	F-0VARC1001496	0. 90
	OVARC1001506	F-0VARC1001506	0. 59
	OVARC1001525	F-0VARC1001525	0. 87
	OVARC1001555	F-0VARC1001555	0. 67
50	OVARC1001577	F-0VARC1001577	0. 94
	0VARC1001610	F-0VARC1001610	0. 38
	0VARC1001611	F-0VARC1001611	0. 34
	0VARC1001702	F-0VARC1001702	0. 88
55	OVARC1001703	F-0VARC1001703	0. 76

	OVARC1001711	F-0VARC1001711	0. 82
	OVARC1001713	F-0VARC1001711	
	OVARC1001713		0.82
5		F-0VARC1001731	0.94
3	OVARC1001762	F-0VARC1001762_	
	OVARC1001766	F-0VARC1001766	0. 94
	OVARC1001767	F-0VARC1001767	0. 70
	OVARC1001768	F-0VARC1001768	0. 32
10	OVARC1001791	F-0VARC1001791	0.74
	OVARC1001802	F-0VARC1001802	0. 76
	OVARC1001809	F-0VARC1001809	0. 63
	OVARC1001820	F-0VARC1001820	0. 94
15	OVARC1001828	F-0VARC1001828	0. 78
	OVARC1001861	F-0VARC1001861	0.94
	OVARC1001873	F-0VARC1001873;	0.31
	OVARC1001879	F-0VARC1001879	0.94
20	OVARC1001880	F-0VARC1001880	0.46
	OVARC1001916	F-0VARC1001916	0.82
	0VARC1001942	F-0VARC1001942	0.72
	OVARC1001943	F-0VARC1001943	0.49
	0VARC1001949	F-0VARC1001949	0.46
25	0VARC1001950	F-0VARC1001950	0.94
	OVARC1001987-	F-0VARC1001987	0. 91
	0VARC1002050	F-0VARC1002050	0. 85
	0VARC1002082	F-0VARC1002082	0.94
30	0VARC1002107	F-0VARC1002107	0.33
	0VARC1002112	F-0VARC1002112	0.94
	OVARC1002127	F-0VARC1002127	0.80
	0VARC1002138	F-0VARC1002138	0.46
35	OVARC1002156	F-0VARC1002156	0.43
	0VARC1002158	F-0VARC1002158	0.94
	0VARC1002165	F-0VARC1002165	0.88
	0VARC1002182	F-0VARC1002182	0.94
40	PLACE1000004	F-PLACE1000004	0.43
	PLACE1000005	F-PLACE1000005	0.37
	PLACE1000007	F-PLACE1000007	0. 72
	PLACE1000014	F-PLACE1000014	0.38
	PLACE1000048	F-PLACE1000048	0.36
45	PLACE1000050	F-PLACE1000050	0. 94
	PLACE1000061	F-PLACE1000061	0.56
	PLACE1000066	F-PLACE1000066	0. 94
	PLACE1000081	F-PLACE1000081	0.37
50	PLACE1000133	F-PLACE1000133	0. 53
	PLACE1000142	F-PLACE1000142	0.94
	PLACE1000184	F-PLACE1000184	0. 73
	PLACE1000185	F-PLACE1000185	0. 86
55	PLACE1000213	F-PLACE1000213	0.34

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	PLACE1000236	F-PLACE1000236	0. 76
	PLACE1000246	F-PLACE1000246	0.91
	PLACE1000308	F-PLACE1000308	0. 40
5	PLACE1000347	F-PLACE1000347_	0. 48
	PLACE1000374	F-PLACE1000374	0. 58
	PLACE1000380	F-PLACE1000380	0. 94
	PLACE1000383	F-PLACE1000383	0.39
10	PLACE1000401	F-PLACE1000401	0. 94
	PLACE1000406	F-PLACE1000406	0.94
	PLACE1000420	F-PLACE1000420	0.82
	PLACE1000435	F-PLACE1000435	0. 32
15	PLACE1000444	F-PLACE1000444	0.41
	PLACE1000453	F-PLACE1000453	0.76
	PLACE1000492	F-PLACE1000492	0.65
	PLACE1000547	F-PLACE1000547	0.94
20	PLACE1000562	F-PLACE1000562	0.94
	PLACE1000564	F-PLACE1000564	0.45
	PLACE1000588	F-PLACE1000588	0.67
	PLACE1000596	F-PLACE1000596	0.60
25	PLACE1000610	F-PLACE1000610	0.94
25	PLACE1000611	F-PLACE1000611	0.48
	PLACE1000636	F-PLACE1000636	0.92
	PLACE1000653	F-PLACE1000653	0. 57
	PLACE1000656	F-PLACE1000656	0.94
30	PLACE1000706	F-PLACE1000706	0.94
	PLACE1000712	F-PLACE1000712	0. 78
	PLACE1000716	F-PLACE1000716	0. 94
	PLACE1000748	F-PLACE1000748	0. 52
35	PLACE1000755	F-PLACE1000755	0.87
	PLACE1000769	F-PLACE1000769	0. 94
	PLACE1000785	F-PLACE1000785	0. 33
	PLACE1000786	F-PLACE1000786	0.88
40	PLACE1000793	F-PLACE1000793	0. 79
	PLACE1000798	F-PLACE1000798	0.59
	PLACE1000849	F-PLACE1000849	0. 64
	PLACE1000856	F-PLACE1000856	0.40
45	PLACE1000863	F-PLACE1000863	0.94
	PLACE1000909	F-PLACE1000909	0.94
	PLACE1000948	F-PLACE1000948	0.92
	PLACE1000972	F-PLACE1000972	0.73
50	PLACE1000977	F-PLACE1000977	0.72
	PLACE1000987	F-PLACE1000987	0.93
	PLACE1001000		0.37
	PLACE1001010		0.50
55	PLACE1001024		0.94
	PLACE1001036	F-PLACE TOUTUS6	0. 51

	PLACE1001054	F-PLACE1001054	0.88
	PLACE1001062	F-PLACE1001062	0.57
	PLACE1001092	F-PLACE1001092	0.94
5	PLACE1001104	F-PLACE1001104	0.38
	PLACE1001168	F-PLACE1001168	0.41
	PLACE1001171	F-PLACE1001171	0.36
	PLACE1001185	F-PLACE1001185	0.93
10	PLACE1001238	F-PLACE1001238	0.87
	PLACE1001241	F-PLACE1001241	0. 67
	PLACE1001257	F-PLACE1001257	0.40
	PLACE1001280	F-PLACE1001280	0.70
15	PLACE1001294	F-PLACE1001294	0.49
	PLACE1001304	F-PLACE1001304	0.94
	PLACE1001311	F-PLACE1001311,	0.53
	PLACE1001351	F-PLACE1001351	0.86
22	PLACE1001366	F-PLACE1001366	0.39
20	PLACE1001383	F-PLACE1001383	0.94
	PLACE1001387	F-PLACE1001387	0. 34
	PLACE1001399	F-PLACE1001399	0.38
	PLACE1001412	F-PLACE1001412	0.94
25	PLACE1001456	F-PLACE1001456	0.42
	PLACE1001484	F-PLACE1001484	0.88
	PLACE1001503	F-PLACE1001503	0.85
	PLACE1001545	F-PLACE1001545	0. 92
30	PLACE1001551	F-PLACE1001551	0.39
	PLACE1001570	F-PLACE1001570	0. 40
	PLACE1001602	F-PLACE1001602	0.60
	PLACE1001608	F-PLACE1001608	0.40
35	PLACE1001610	F-PLACE1001610	0. 43
	PLACE1001632	F-PLACE1001632	0.48
	PLACE1001634	F-PLACE1001634	0.60
	PLACE1001692	F-PLACE1001692	0.74
40	PLACE1001729	F-PLACE1001729	0.69
	PLACE1001739	F-PLACE1001739	0.49
	PLACE1001740	F-PLACE1001740	0.50
	PLACE1001761	F-PLACE1001761	0.71
45	PLACE1001771		
	PLACE1001781	F-PLACE1001781	0. 57
	PLACE1001810	F-PLACE1001810	0.57
	PLACE1001817	F-PLACE1001817	0.91
50	PLACE1001844	F-PLACE1001844	0.34
	PLACE1001869	F-PLACE1001869	0.34
	PLACE1001912	F-PLACE1001912	0.82
	PLACE1001920	F-PLACE1001920	0.85
	PLACE1001928	F-PLACE1001928	0.60
55	PLACE1001983	F-PLACE1001983	0.39

	PLACE1001989	F-PLACE1001989	0.:74
	PLACE1002004	F-PLACE1002004	0.70
	PLACE1002046		0. 81
5	PLACE1002072	F-PLACE1002072	
•	PLACE1002073	F-PLACE1002073	0.70
	PLACE1002090	F-PLACE1002090	0. 64
	PLACE1002030	F-PLACE1002115	0.60
10	PLACE 1002119	F-PLACE1002119	0.71
10		F-PLACE1002140	0. 71
	PLACE1002140	F-PLACE1002140	0. 56
	PLACE1002163		
	PLACE1002170	F-PLACE1002170	0.60
15	PLACE1002171	F-PLACE1002171	0.72
	PLACE1002213	F-PLACE1002213	0.35
	PLACE1002342	F-PLACE1002342,	
	PLACE1002395		0.87
20	PLACE1002433	F-PLACE1002433	0. 73
	PLACE1002437	F-PLACE1002437	0. 46
	PLACE1002438	F-PLACE1002438	0.41
	PLACE1002450	F-PLACE1002450	0.48
25	PLACE1002465	F-PLACE1002465	0.34
	PLACE1002474	F-PLACE1002474	0.89
	PLACE1002499	F-PLACE1002499	0.94
	PLACE1002529	F-PLACE1002529	0. 45
	PLACE1002532	F-PLACE1002532	0. 76
30	PLACE1002571	F-PLACE1002571	0.40
	PLACE1002625	F-PLACE1002625	0.36
	PLACE1002655	F-PLACE1002655	0.41
	PLACE1002665	F-PLACE1002665	0.80
35	PLACE1002685	F-PLACE1002685	0. 31
	PLACE1002714	F-PLACE1002714	0.90
	PLACE1002722	F-PLACE1002722	0. 43
	PLACE1002775	F-PLACE1002775	0.94
40	PLACE1002794	F-PLACE1002794	0.90
	PLACE1002811	F-PLACE1002811	0.88
	PLACE1002815	F-PLACE1002815	0.40
	PLACE1002816	F-PLACE1002816	0. 73
45	PLACE1002834		0. 41
	PLACE1002839	F-PLACE1002839	0. 73
	PLACE1002851	F-PLACE1002851	0.90
	PLACE1002908	F-PLACE1002908	0.94
50	PLACE1002941	F-PLACE1002941	0. 33
50	PLACE1002991	F-PLACE1002991	0.39
	PLACE1002993	F-PLACE1002993	0.94
	PLACE1002996	F-PLACE1002996	0.40
	PLACE1003025		0.77
55	PLACE1003027	F-PLACE1003027	0.54

	PLACE1003044	F-PLACE1003044	0. 71
	PLACE1003045	F-PLACE1003045	0. 89
	PLACE1003092	F-PLACE1003092	0. 59
5	PLACE1003100	F-PLACE1003100	0, 38
	PLACE1003108	F-PLACE1003108	0. 94
	PLACE1003136	F-PLACE1003136	0.87
	PLACE1003145	F-PLACE1003145	0.94
10	PLACE1003174	F-PLACE1003174	0.46
	PLACE1003176	F-PLACE1003176	0.48
	PLACE1003190	F-PLACE1003190	0.57
	PLACE1003200	F-PLACE1003200	0.47
15	PLACE1003249	F-PLACE1003249	0.33
-	PLACE1003256	F-PLACE1003256	0.57
	PLACE1003296	F-PLACE1003296,	0.41
	PLACE1003302	F-PLACE1003302	0.94
20	PLACE1003334	F-PLACE1003334	0.78
20	PLACE1003342	F-PLACE1003342	0.34
	PLACE1003353	F-PLACE1003353	0.71
	PLACE1003366	F-PLACE1003366	0.66
	PLACE1003369	F-PLACE1003369	0. 91
25	PLACE1003383	F-PLACE1003383	0. 70
	PLACE1003394	F-PLACE1003394	0. 94
	PLACE1003454	F-PLACE1003454	0.94
	PLACE1003493	F-PLACE1003493	0. 93
30	PLACE1003537	F-PLACE1003537	0. 94
	PLACE1003553	F-PLACE1003553	0.34
	PLACE1003592	F-PLACE1003592	0. 33
	PLACE1003596	F-PLACE1003596	0.89
35	PLACE1003602	F-PLACE1003602	0. 94
	PLACE1003605	F-PLACE1003605	0.94
	PLACE1003611	F-PLACE1003611	0.80
	PLACE1003625	F-PLACE1003625	0.39
40	PLACE1003669	F-PLACE1003669	0. 79 0. 43
	PLACE1003704 PLACE1003709	F-PLACE1003704 F-PLACE1003709	0. 43
	PLACE 1003709	F-PLACE1003703	0. 74
	PLACE1003711	F-PLACE1003711	0. 63
45		F-PLACE1003723	0. 03
	PLACE1003738 PLACE1003762	F-PLACE1003762	0.94
	PLACE1003702 PLACE1003771	F-PLACE1003702	0. 56
	PLACE1003771	F-PLACE1003771	0.39
50	PLACE1003784	F-PLACE1003784	0.61
	PLACE1003795	F-PLACE1003795	0.32
	PLACE1003793	F-PLACE1003870	0. 94
	PLACE1003876	F-PLACE1003885	0. 94
55	PLACE1003886	F-PLACE1003886	0. 94

	PLACE1003888	F-PLACE1003888	0. 36
	PLACE1003892	F-PLACE1003892	0. 92
	PLACE1003903	F-PLACE1003903	0. 68
5	PLACE1003915	F-PLACE1003915	0. 54
	PLACE1003923	F-PLACE1003923	0. 52
	PLACE1003936	F-PLACE1003936	0. 67
	PLACE1003968	F-PLACE1003968	0.59
10	PLACE1004103	F-PLACE1004103	0.94
	PLACE1004104	F-PLACE1004104	0. 44
	PLACE1004114	F-PLACE1004114	0. 91
	PLACE1004128	F-PLACE1004128	0.72
15	PLACE1004149	F-PLACE1004149	0. 41
	PLACE1004156	F-PLACE1004156	0. 50
	PLACE1004161	F-PLACE1004161,	0.77
	PLACE1004183	F-PLACE1004183	0.62
20	PLACE1004197	F-PLACE1004197	0.39
	PLACE1004203	F-PLACE1004203	0.94
	PLACE1004258	F-PLACE1004258	0.46
	PLACE1004270	F-PLACE1004270	0.65
25	PLACE1004277	F-PLACE1004277	0.86
25	PLACE1004289	F-PLACE1004289	0.79
	PLACE1004302	F-PLACE1004302	0.57
	PLACE1004316	F-PLACE1004316	0. 64
00	PLACE1004358	F-PLACE1004358	0.85
30	PLACE1004376	F-PLACE1004376	0.64
	PLACE1004388	F-PLACE1004388	0.94
	PLACE1004405	F-PLACE1004405	0.82
	PLACE1004428	F-PLACE1004428	0. 68
35 .	PLACE1004437	F-PLACE1004437	0. 94
	PLACE1004451	F-PLACE1004451	0. 39
	PLACE1004460	F-PLACE1004460	0. 92
	PLACE1004473	F-PLACE1004473	0. 94
40	PLACE1004510	F-PLACE1004510	0. 94
	PLACE1004516	F-PLACE1004516	0. 58
	PLACE1004548 PLACE1004550	F-PLACE1004548 F-PLACE1004550	0.59
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45	PLACE1004504 PLACE1004629	F-PLACE1004584	
	PLACE1004645	F-PLACE1004645	0. 34 0. 59
	PLACE1004646	F-PLACE1004646	
	PLACE1004646	F-PLACE1004644	0. 35 0. 31
50	PLACE1004672	F-PLACE1004672	0. 83
	PLACE1004674	F-PLACE1004674	0. 90
	PLACE1004686	F-PLACE1004686	0. 30
·	PLACE1004691	F-PLACE1004691	0. 89
55	PLACE1004722	F-PLACE1004722	0. 48

	PLACE1004736	F-PLACE1004736	0. 80
	PLACE1004740	F-PLACE1004740	0.34
	PLACE1004743	F-PLACE1004743	0. 70
5	PLACE 1004751	F-PLACE1004751	0. 49
	PLACE1004777	F-PLACE1004777	0. 72
	PLACE1004804	F-PLACE1004804	0.46
	PLACE1004814	F-PLACE1004814	0. 70
10	PLACE1004824	F-PLACE1004824	0. 63
	PLACE1004868	F-PLACE1004868	0.94
	PLACE1004885	F-PLACE1004885	0. 63
	PLACE1004902	F-PLACE1004902	0. 56
15	PLACE1004918	F-PLACE1004918	0. 85
	PLACE1004930	F-PLACE1004930	0. 83
	PLACE1004937	F-PLACE1004937;	0. 46
	PLACE1004969	F-PLACE1004969	0.62
20	PLACE1004982	F-PLACE1004982	0.61
	PLACE1005026	F-PLACE1005026	0.81
	PLACE1005027	F-PLACE1005027	0.91
	PLACE1005046	F-PLACE1005046	0.31
25	PLACE1005055	F-PLACE1005055	0. 57
25	PLACE1005066	F-PLACE1005066	0. 68
	PLACE1005101	F-PLACE1005101	0.94
	PLACE1005102	F-PLACE1005102	0. 94
	PLACE1005181	F-PLACE1005181	0. 94
30	PLACE1005187	F-PLACE1005187	0. 94
	PLACE1005206	F-PLACE1005206	0.67
	PLACE1005232	F-PLACE1005232	0. 72
	PLACE1005243	F-PLACE1005243	0. 81
35	PLACE1005261	F-PLACE1005261	0. 75
	PLACE1005266	F-PLACE1005266	0. 55
	PLACE1005277	F-PLACE1005277	0.43
	PLACE1005287	F-PLACE1005287	0.77
40	PLACE1005305	F-PLACE1005305	0.94
	PLACE1005308	F-PLACE1005308	0.46
	PLACE1005313	F-PLACE1005313	0.94
	PLACE1005327	F-PLACE1005327	0.82
45	PLACE1005331	F-PLACE1005331	0.94
	PLACE1005335 PLACE1005373	F-PLACE1005335	0. 94 0. 71
	PLACE1005373 PLACE1005374	F-PLACE1005373 F-PLACE1005374	0. 71
	PLACE1005374 PLACE1005480	F-PLACE1005480	0. 82
50	PLACE1005480	F-PLACE1005481	0. 42
	PLACE1005494	F-PLACE1005494	0. 42
	PLACE1005494 PLACE1005530	F-PLACE1005530	0. 94
	PLACE1005550		0. 86
55	PLACE1005554		0. 34
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	PLACE1005557	F-PLACE1005557	0.80
	PLACE1005595	F-PLACE1005595	0. 69
	PLACE1005603	F-PLACE1005603	0. 92
5	PLACE1005623	F-PLACE1005623	0.68
	PLACE1005630	F-PLACE1005630	0. 58
	PLACE1005646	F-PLACE1005646	0.31
	PLACE1005656	F-PLACE1005656	0.85
10	PLACE1005698	F-PLACE1005698	0. 91
	PLACE1005727	F-PLACE1005727	0.94
	PLACE1005739	F-PLACE1005739	0. 87
	PLACE1005755	F-PLACE1005755	0. 32
15	PLACE1005763	F-PLACE1005763	0.74
	PLACE1005799	F-PLACE1005799	0.47
	PLACE1005803	F-PLACE1005803,	0.88
	PLACE1005804	F-PLACE1005804	0.36
20	PLACE1005813	F-PLACE1005813	0.76
	PLACE1005828	F-PLACE1005828	0.43
	PLACE1005834	F-PLACE1005834	0.89
	PLACE1005851	F-PLACE1005851	0.39
05	PLACE1005876	F-PLACE1005876	0.51
25	PLACE1005890	F-PLACE1005890	0.81
	PLACE1005921	F-PLACE1005921	0.90
	PLACE1005923	F-PLACE1005923	0.70
	PLACE1005925	F-PLACE1005925	0.55
30	PLACE1005934	F-PLACE1005934	0. 94
	PLACE1005936	F-PLACE1005936	0. 41
	PLACE1005951	F-PLACE1005951	0.34
	PLACE1005953	F-PLACE1005953	0. 31
35	PLACE1005955	F-PLACE1005955	0.89
	PLACE1005966	F-PLACE1005966	0.73
	PLACE1005990	F-PLACE1005990	0. 37
	PLACE1006011	F-PLACE1006011	0. 94
40	PLACE1006037	F-PLACE1006037	0. 65
	PLACE1006040	F-PLACE1006040	0.94
	PLACE1006119	F-PLACE1006119	0.52
	PLACE1006139	F-PLACE1006139	0.94
45	PLACE1006157		0.34
	PLACE1006159	F-PLACE1006159	0.50
	PLACE1006167		0.94
	PLACE1006170	F-PLACE1006170	0.94
50	PLACE1006195	F-PLACE1006195	0.31
	PLACE1006196	F-PLACE1006196	0.84
	PLACE1006225	F-PLACE1006225	0.87
	PLACE1006236	F-PLACE1006236	0.92
55	PLACE1006239		0. 53
55	PLACE1006246	F-PLACE1006246	0. 44

	PLACE1006248	F-PLACE1006248	0.49
	PLACE1006325	F-PLACE1006325	0.74
	PLACE1006335	F-PLACE1006335	0. 82
5	PLACE1006357	F-PLACE1006357	0. 33
	PLACE1006385	F-PLACE1006385	0. 54
	PLACE1006412	F-PLACE1006412	0. 79
	PLACE1006414	F-PLACE1006414	0. 52
10	PLACE1006438	F-PLACE1006438	0.94
	PLACE1006445	F-PLACE1006445	0. 37
	PLACE1006469	F-PLACE1006469	0.85
	PLACE1006482	F-PLACE1006482	0. 94
15	PLACE1006488	F-PLACE1006488	0. 94
	PLACE1006492	F-PLACE1006492	0.32
	PLACE1006531	F-PLACE1006531	0. 45
	PLACE1006552	F-PLACE1006552	0.42
20	PLACE1006598	F-PLACE1006598	0.33
	PLACE1006615	F-PLACE1006615	0. 89
	PLACE1006626	F-PLACE1006626	0.69
	PLACE1006673	F-PLACE1006673	0.61
a.e.	PLACE1006678	F-PLACE1006678	0. 53
25	PLACE1006704		0. 47
	PLACE1006731	F-PLACE1006731	0.72
	PLACE1006782	F-PLACE1006782	0. 68
	PLACE1006819	F-PLACE1006819	0. 57
30	PLACE1006829	F-PLACE1006829	0. 47
	PLACE1006883	F-PLACE1006883	0. 88
	PLACE1006901	F-PLACE1006901	0. 43
	PLACE1006917	F-PLACE1006917	0.56
35	PLACE1006932	F-PLACE1006932	0.45
	PLACE1006935	F-PLACE1006935	0.58
	PLACE1006956	F-PLACE1006956	0.77
	PLACE1006958 PLACE1006961	F-PLACE1006958 F-PLACE1006961	0. 75 0. 34
40	PLACE1006961	F-PLACE1006961	0. 94
	PLACE1006962 PLACE1006966	F-PLACE1006962	0. 56
	PLACE1006988	F-PLACE1006989	0. 47
	PLACE1000989		0. 47
45	PLACE1007014	F-PLACE1007021	0. 74
	PLACE1007053	F-PLACE1007053	0.73
	PLACE1007068	F-PLACE1007068	0. 43
	PLACE1007105	F-PLACE1007105	0. 88
50	PLACE1007112	F-PLACE1007112	0. 82
	PLACE1007178	F-PLACE1007178	0. 91
	PLACE1007226	F-PLACE1007226	0. 79
	PLACE1007238	F-PLACE1007238	0.94
55	PLACE1007239	F-PLACE1007239	0.94

	· PLACE1007242	F-PLACE1007242	0.48
	PLACE1007243	F-PLACE1007243	0. 77
•	PLACE1007257	F-PLACE1007257	0. 54
5	PLACE1007274	F-PLACE1007274	0. 94
	PLACE1007282	F-PLACE1007282	0. 94
	PLACE1007301	F-PLACE1007301	0.34
	PLACE1007317	F-PLACE1007317	0. 43
10	PLACE1007342	F-PLACE1007342	0.72
	PLACE1007346	F-PLACE1007346	0.92
	PLACE1007367	F-PLACE1007367	0.43
	PLACE1007375	F-PLACE1007375	0. 85
15	PLACE1007386	F-PLACE1007386	0. 58
· ·	PLACE1007402	F-PLACE1007402	0.74
	PLACE1007409	F-PLACE1007409,	0.39
	PLACE1007416	F-PLACE1007416	0.80
20	PLACE1007450	F-PLACE1007450	0.41
20	PLACE1007452	F-PLACE1007452	0.68
	PLACE1007454	F-PLACE1007454	0.39
	PLACE1007460	F-PLACE1007460	0.58
	PLACE1007484	F-PLACE1007484	0.94
25	PLACE1007488	F-PLACE1007488	0.50
	PLACE1007507	F-PLACE1007507	0.44
	PLACE1007511	F-PLACE1007511	0.86
	PLACE1007524	F-PLACE1007524	0.66
30	PLACE1007537	F-PLACE1007537	0.93
	PLACE1007544	F-PLACE1007544	0.94
	PLACE1007547	F-PLACE1007547	0.65
	PLACE1007583	F-PLACE1007583	0.87
35	PLACE1007598	F-PLACE1007598	0.39
	PLACE1007618	F-PLACE1007618	0.62
	PLACE1007621	F-PLACE1007621	0.63
	PLACE1007632	F-PLACE1007632	0.75
40	PLACE1007645	F-PLACE1007645	0.94
	PLACE1007649	F-PLACE1007649	0. 56
	PLACE1007688	F-PLACE1007688	0.94
	PLACE1007690	F-PLACE1007690	0. 44
45	PLACE1007697	F-PLACE1007697	0.93
	PLACE1007705	F-PLACE1007705	0.61
	PLACE1007706	F-PLACE1007706	0.47
	PLACE1007725	F-PLACE1007725	0.61
50	PLACE1007729	F-PLACE1007729	0.53
<del></del>	PLACE1007730	F-PLACE1007730	0.31
	PLACE1007746	F-PLACE1007746	0.81
	PLACE1007791	F-PLACE1007791	0. 46 0. 73
	PLACE1007810		0. 73
55	PLACE1007843	F-FLAUE 100/843	0.31

	PLACE1007846	F-PLACE1007846	0.38
	PLACE1007858	F-PLACE1007858	0.79
	PLACE1007897	F-PLACE1007897	0.94
5	PLACE1007908	F-PLACE1007908	0.94
	PLACE1007946	F-PLACE1007946	0.52
	PLACE1007954	F-PLACE1007954	0.43
	PLACE1007955	F-PLACE1007955	0.41
10	PLACE1007958	F-PLACE1007958	0.37
	PLACE1007969	F-PLACE1007969	0.87
	PLACE1007990	F-PLACE1007990	0.94
	PLACE1008000	F-PLACE1008000	0.94
15	PLACE1008002	F-PLACE1008002	0.93
	PLACE1008044	F-PLACE1008044	0.72
	PLACE1008095	F-PLACE1008095,	0.94
	PLACE1008122	F-PLACE1008122	0. 94
20	PLACE1008129	F-PLACE1008129	0.80
20	PLACE1008132	F-PLACE1008132	0.37
	PLACE1008177	F-PLACE1008177	0.74
	PLACE1008198	F-PLACE1008198	0.94
	PLACE1008209	F-PLACE1008209	0. 54
25	PLACE1008273	F-PLACE1008273	0.94
	PLACE1008275	F-PLACE1008275	0. 82
	PLACE1008280	F-PLACE1008280	0.34
	PLACE1008309	F-PLACE1008309	0. 90
30	PLACE1008329	F-PLACE1008329	0. 38
	PLACE1008356	F-PLACE1008356	0. 93
	PLACE1008368	F-PLACE1008368	0. 43
	PLACE1008398	F-PLACE1008398	0. 78
35	PLACE1008401	F-PLACE1008401	0.81
	PLACE1008402	F-PLACE1008402	0. 88
	PLACE1008429	F-PLACE1008429	0. 39
	PLACE1008457	F-PLACE1008457	0. 77
40	PLACE1008465	F-PLACE1008465	0. 94
	PLACE1008488	F-PLACE1008488	0. 42
	PLACE1008524	F-PLACE1008524	0.41
	PLACE1008531	F-PLACE1008531	0.94
45	PLACE1008532	F-PLACE1008532	0.61
	PLACE1008533	F-PLACE1008533	0. 45
	PLACE1008568	F-PLACE1008568	0.65
	PLACE1008603	F-PLACE1008603	0. 42
50	PLACE1008626	F-PLACE1008626	0. 94
	PLACE1008627	F-PLACE1008627	0. 72
	PLACE1008629	F-PLACE1008629	0.33
	PLACE1008650	F-PLACE1008650 F-PLACE1008693	0.79
22	PLACE1008693		0. 32 0. 39
55	PLACE1008696	F-PLACE1008696	U. 39

	PLACE1008790	F-PLACE1008790	0.81
	PLACE1008808	F-PLACE1008808	0.51
	PLACE1008813	F-PLACE1008813	0.36
5	PLACE1008854	F-PLACE1008854	0.81
	PLACE1008867	F-PLACE1008867	0.34
,	PLACE1008887	F-PLACE1008887	0.49
	PLACE1008902	F-PLACE1008902	0.91
10	PLACE1008925	F-PLACE1008925	0.94
	PLACE1008934	F-PLACE1008934	0.70
	PLACE1009027	F-PLACE1009027	0.32
	PLACE1009045	F-PLACE1009045	0.42
15	PLACE1009060	F-PLACE1009060	0.69
	PLACE1009090	F-PLACE1009090	0.46
	PLACE1009091	F-PLACE1009091,	0.90
	PLACE1009094	F-PLACE1009094	0.44
20	PLACE1009099	F-PLACE1009099	0.56
	PLACE1009110	F-PLACE1009110	0.84
	PLACE1009111	F-PLACE1009111	0.80
	PLACE1009113	F-PLACE1009113	0.88
25	PLACE1009130	F-PLACE1009130	0.60
25	PLACE1009158	F-PLACE1009158	0.41
	PLACE1009166	F-PLACE1009166	0.94
	PLACE1009174	F-PLACE1009174	0.38
22	PLACE1009186	F-PLACE1009186	0. 59
30	PLACE1009190	F-PLACE1009190	0.66
	PLACE1009230	F-PLACE1009230	0.87
	PLACE1009298	F-PLACE1009298	0.41
	PLACE1009319	F-PLACE1009319	0. 53
35	PLACE1009328	F-PLACE1009328	0.66
	PLACE1009335	F-PLACE1009335	0. 47
	PLACE1009338	F-PLACE1009338	0.70
	PLACE1009368	F-PLACE1009368	0.94
40	PLACE1009375	F-PLACE1009375	0.76
	PLACE1009388	F-PLACE1009388	0.46
	PLACE1009404 PLACE1009434	F-PLACE1009404 F-PLACE1009434	0. 72 0. 50
	PLACE1009434 PLACE1009443	F-PLACE1009434	0. 90
45	PLACE1009444	F-PLACE1009444	0.33
	PLACE1009459	F-PLACE1009459	0. 94
	PLACE1009458	F-PLACE1009468	0.88
	PLACE1009476	F-PLACE1009476	0.84
50	PLACE1009524	F-PLACE1009524	0.88
	PLACE1009542	F-PLACE1009542	0.55
	PLACE1009571	F-PLACE1009571	0.61
	PLACE1009581	F-PLACE1009581	0.94
55	PLACE1009596	F-PLACE1009596	0. 44

	PLACE1009607	F-PLACE1009607	0. 47
	PLACE1009622	F-PLACE1009622	0.65
	PLACE1009659	F-PLACE1009659	0.36
5	PLACE1009665	F-PLACE1009665	0.43
	PLACE1009670	F-PLACE1009670	0.33
	PLACE1009708	F-PLACE1009708	0.80
	PLACE1009721	F-PLACE1009721	0.81
10	PLACE1009731	F-PLACE1009731	0.47
	PLACE1009763	F-PLACE1009763	0.94
	PLACE1009794	F-PLACE1009794	0.33
	PLACE1009798	F-PLACE1009798	0.34
15	PLACE1009845	F-PLACE1009845	0.66
	PLACE1009861	F-PLACE1009861	0.72
	PLACE1009886	F-PLACE1009886,	0.63
	PLACE1009908	F-PLACE1009908	0.92
00	PLACE1009921	F-PLACE1009921	0.70
20	- PLACE1009971	F-PLACE1009971	0.40
	PLACE1009992	F-PLACE1009992	0.89
	PLACE1009995	F-PLACE1009995	0.88
	PLACE1009997	F-PLACE1009997	0.41
25	PLACE1010023	F-PLACE1010023	0.90
	PLACE1010031	F-PLACE1010031	0.71
	PLACE1010053	F-PLACE1010053	0.51
	PLACE1010074	F-PLACE1010074	0. 75
30	PLACE1010076	F-PLACE1010076	0. 53
	PLACE1010096	F-PLACE1010096	0. 44
	PLACE1010102	F-PLACE1010102	0. 77
	PLACE1010105	F-PLACE1010105	0. 58
35	PLACE1010106	F-PLACE1010106	0. 39
	PLACE1010134	F-PLACE1010134	0.85
	PLACE1010148	F-PLACE1010148	0. 32
	PLACE1010152	F-PLACE1010152	0.94
40	PLACE1010194	F-PLACE1010194	0.94
	PLACE1010202	F-PLACE1010202	0. 33
	PLACE1010261	F-PLACE1010261	0.36
	PLACE1010274	F-PLACE1010274	0.94
45	PLACE1010293	F-PLACE1010293	0.55
	PLACE1010310	F-PLACE1010310	0. 41 0. 40
	PLACE1010321	F-PLACE1010321	
	PLACE1010324 PLACE1010329	F-PLACE1010324 F-PLACE1010329	0. 40 0. 36
50		F-PLACE1010329	0.36
	PLACE1010362	F-PLACE1010362	0. 34
	PLACE1010364 PLACE1010383	F-PLACE1010384	0. 37
	PLACE1010383 PLACE1010401	F-PLACE1010363	0. 53
**	PLACE1010401	F-PLACE1010481	0. 74
55	PENUE IU IU481	1-50001010401	0. /4

	PLACE1010491	F-PLACE1010491	0.31
	PLACE1010492	F-PLACE1010492	0.39
	PLACE1010522	F-PLACE1010522	0.92
5	PLACE1010529	F-PLACE1010529	0.65
	PLACE1010547	F-PLACE1010547	0. 82
	PLACE1010580	F-PLACE1010580	0.72.
	PLACE1010599	F-PLACE1010599	0.83
10	PLACE1010616	F-PLACE1010616	0. 54
	PLACE1010622	F-PLACE1010622	0. 92
	PLACE1010629	F-PLACE1010629	0.94
	PLACE1010630	F-PLACE1010630	0.37
15	PLACE1010661	F-PLACE1010661	0. 56
	PLACE1010714	F-PLACE1010714	0. 68
	PLACE1010720	F-PLACE1010720,	0.90
	PLACE1010743	F-PLACE1010743	0. 75
20	PLACE1010761	F-PLACE1010761	0.69
20	PLACE1010771	F-PLACE1010771	0.94
	PLACE1010786	F-PLACE1010786	0.80
	PLACE1010800	F-PLACE1010800	0. 41
	PLACE1010811	F-PLACE1010811	0.94
25	PLACE1010833	F-PLACE1010833	0. 94
	PLACE1010856	F-PLACE1010856	0. 37
	PLACE1010857	F-PLACE1010857	0. 54
	PLACE1010870	F-PLACE1010870	0.50
30	PLACE1010877	F-PLACE1010877	0. 77
	PLACE1010900	F-PLACE1010900	0.70
	PLACE1010917	F-PLACE1010917	0.94
	PLACE1010925	F-PLACE1010925	0. 79
35	PLACE1010926	F-PLACE1010926	0.94
	PLACE1010942	F-PLACE1010942 F-PLACE1010944	0. 36 0. 61
	PLACE1010944	F-PLACE1010944	0. 83
	PLACE1010954 PLACE1010960	F-PLACE1010960	0. 38
40	PLACE1010965	F-PLACE1010965	0. 68
	PLACE1010905	F-PLACE1011026	0. 38
	PLACE1011041	F-PLACE1011041	0. 94
	PLACE1011046	F-PLACE1011046	0.85
45	PLACE1011054	F-PLACE1011054	0. 65
•	PLACE1011056	F-PLACE1011056	0.82
	PLACE1011090	F-PLACE1011090	0.94
	PLACE1011114	F-PLACE1011114	0. 94
50	PLACE1011160	F-PLACE1011160	0.62
	PLACE1011214	F-PLACE1011214	0.40
	PLACE1011219	F-PLACE1011219	0.48
	PLACE1011221	F-PLACE1011221	0.94
55	PLACE1011229	F-PLACE1011229	0.46

	PLACE1011263	F-PLACE1011263	0. 94
	PLACE1011273	F-PLACE1011273	0. 32
	PLACE1011291	F-PLACE1011291	0. 55
5	PLACE1011310	F-PLACE1011310	0.74
	PLACE1011325	F-PLACE1011325	0.39
	PLACE1011332	F-PLACE1011332	0. 74
	PLACE1011371	F-PLACE1011371	0. 32
10	PLACE1011399	F-PLACE1011399	0. 89
	PLACE1011433	F-PLACE1011433	0. 33
	PLACE1011472	F-PLACE1011472	0. 59
	PLACE1011477	F-PLACE1011477	0. 80
15	PLACE1011563	F-PLACE1011563	0.83
,,	PLACE1011586	F-PLACE1011586	0. 82
	PLACE1011635	F-PLACE1011635,	
	PLACE1011646	F-PLACE1011646	0. 37
	PLACE1011664	F-PLACE1011664	0. 65
20	PLACE1011675	F-PLACE1011675	0.89
	PLACE1011682	F-PLACE1011682	0.81
	PLACE1011725	F-PLACE1011725	0.86
	PLACE1011783	F-PLACE1011783	0. 78
25	PLACE1011858	F-PLACE1011858	0.76
	PLACE1011875	F-PLACE1011875	0.49
	PLACE1011891	F-PLACE1011891	0.34
	PLACE1011896	F-PLACE1011896	0.34
30	PLACE1011922	F-PLACE1011922	0.46
	PLACE1011923	F-PLACE1011923	0.74
	PLACE1011982	F-PLACE1011982	0.71
	PLACE1012031	F-PLACE1012031	0.80
35	PLACE2000006	F-PLACE2000006	0.61
	PLACE2000007	F-PLACE2000007	0.37
	PLACE2000014	F-PLACE2000014	0.67
	PLACE2000015	F-PLACE2000015	0.53
40	PLACE2000021	F-PLACE2000021	0.36
	PLACE2000030	F-PLACE2000030	0.83
	PLACE2000033	F-PLACE2000033	0. 44
	PLACE2000034	F-PLACE2000034	0.88
45	PLACE2000039	F-PLACE2000039	0. 67
43	PLACE2000047	F-PLACE2000047	0. 84
	PLACE2000050	F-PLACE2000050	0.41
•	PLACE2000062	F-PLACE2000062	0.94
	PLACE2000072	F-PLACE2000072	0.86
50	PLACE2000097	F-PLACE2000097	0. 94
	PLACE2000100	F-PLACE2000100	0. 75
	PLACE2000111	F-PLACE2000111	0. 62
	PLACE2000124	F-PLACE2000124	0. 77
55	PLACE2000164	F-PLACE2000164	0. 57

	•		
	PLACE2000170	F-PLACE2000170	0.60
	PLACE2000172	F-PLACE2000172	0.55
	PLACE2000223	F-PLACE2000223	0.54
5	PLACE2000235	F-PLACE2000235	0.94
	PLACE2000246	F-PLACE2000246	0.94
	PLACE2000274	F-PLACE2000274	0.82
	PLACE2000302	F-PLACE2000302	0.92
10	PLACE2000341	F-PLACE2000341	0.58
	PLACE2000359	F-PLACE2000359	0.33
	PLACE2000371	F-PLACE2000371	0.71
	PLACE2000373	F-PLACE2000373	0. 65
15	PLACE2000399	F-PLACE2000399	0.41
	PLACE2000404	F-PLACE2000404	0. 94
	PLACE2000411	F-PLACE2000411,	
	PLACE2000427	F-PLACE2000427	0.60
•	PLACE2000433	F-PLACE2000433	0. 61
20	PLACE2000438	F-PLACE2000438	0.56
	PLACE2000458	F-PLACE2000458	0. 82
	PLACE3000009	F-PLACE3000009	0. 56
	PLACE3000020	F-PLACE3000020	0. 94
25	PLACE3000029	F-PLACE3000029	0. 94
	PLACE3000059	F-PLACE3000059	0.51
	PLACE3000070	F-PLACE3000070	0. 67
	PLACE3000103	F-PLACE3000103	0.34
30	PLACE3000121	F-PLACE3000121	0. 32
	PLACE3000142	F-PLACE3000142	0.67
	PLACE3000145	F-PLACE3000145	0.84
	PLACE3000147	F-PLACE3000147	0. 57
35	PLACE3000148	F-PLACE3000148	0.94
	PLACE3000155	F-PLACE3000155	0.42
	PLACE3000156	F-PLACE3000156	0.40
	PLACE3000157	F-PLACE3000157	0. 75
40	PLACE3000160	F-PLACE3000160	0.44
	PLACE3000194	F-PLACE3000194	0. 37
	PLACE3000197	F-PLACE3000197	0.37
	PLACE3000221	F-PLACE3000221	0.49
45	PLACE3000226	F-PLACE3000226	0. 39
45	PLACE3000242	F-PLACE3000242	0.94
	PLACE3000244	F-PLACE3000244	0. 56
	PLACE3000254	F-PLACE3000254	0.60
	PLACE3000304	F-PLACE3000304	0. 42
50	PLACE3000310	F-PLACE3000310	0. 71
	PLACE3000320	F-PLACE3000320	0. 37
	PLACE3000322	F-PLACE3000322	0. 46
	PLACE3000331	F-PLACE3000331	0. 31
55	PLACE3000339	F-PLACE3000339	0. 45

	PLACE3000350	F-PLACE3000350	0. 52 <sup>.</sup>
	PLACE3000352	F-PLACE3000352	0. 65
	PLACE3000353	F-PLACE3000353	0.71
5	PLACE3000363	F-PLACE3000363	0. 45
	PLACE3000405	F-PLACE3000405	0. 52
	PLACE3000413	F-PLACE3000413	0. 85
	PLACE3000416	F-PLACE3000416	0.94
10	PLACE3000425	F-PLACE3000425	0.40
	PLACE3000455	F-PLACE3000455	0.45
	PLACE3000477	F-PLACE3000477	0.79
	PLACE4000009	F-PLACE4000009	0.74
15	PLACE4000014	F-PLACE4000014	0.46
	PLACE4000034	F-PLACE4000034	0.89
	PLACE4000049	F-PLACE4000049,	0.68
	PLACE4000052	F-PLACE4000052	0.80
20	PLACE4000063	F-PLACE4000063	0.69
	PLACE4000089	F-PLACE4000089	0.44
	PLACE4000106	F-PLACE4000106	0.34
	PLACE4000128	F-PLACE4000128	0.89
25	PLACE4000129	F-PLACE4000129	0.61
25	PLACE4000131	F-PLACE4000131	0.55
	PLACE4000147	F-PLACE4000147	0.94
	PLACE4000156	F-PLACE4000156	0.37
	PLACE4000192	F-PLACE4000192	0.88
30	PLACE4000211	F-PLACE4000211	0.64
	PLACE4000222	F-PLACE4000222	0.40
	PLACE4000230	F-PLACE4000230	0. 49
	PLACE4000259	F-PLACE4000259	0.84
35	PLACE4000261	F-PLACE4000261	0.42
	PLACE4000269	F-PLACE4000269	0.60
	PLACE4000270	F-PLACE4000270	0.61
	PLACE4000300	F-PLACE4000300	0.85
40	PLACE4000323	F-PLACE4000323	0.76
	PLACE4000326	F-PLACE4000326	0.86
	PLACE4000369	F-PLACE4000369	0.87
•	PLACE4000387	F-PLACE4000387	0. 41
45	PLACE4000392	F-PLACE4000392	0.39
·	PLACE4000431	F-PLACE4000431	0.88
	PLACE4000445	F-PLACE4000445	0.40
	PLACE4000450 PLACE4000465	F-PLACE4000450 F-PLACE4000465	0. 89 0. 38
50	PLACE4000465	F-PLACE4000465	0. 44
	PLACE4000487	F-PLACE4000489	0. 51
	PLACE4000489	F-PLACE4000489	0.31
	PLACE4000521	F-PLACE4000521	0. 33
55	PLACE4000522	F-PLACE4000558	0. 44
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•	PLACE4000581	F-PLACE4000581	0. 59
	PLACE4000590	F-PLACE4000590	0.38
	PLACE4000593	F-PLACE4000593	0.33
5	PLACE4000612	F-PLACE4000612	0. 37
	PLACE4000638	F-PLACE4000638	0.38
	PLACE4000650	F-PLACE4000650	0.59
	PLACE4000654	F-PLACE4000654	0. 57
10	PLACE4000670	F-PLACE4000670	0.71
	SKNMC1000011	F-SKNMC1000011	0.94
	SKNMC1000013	F-SKNMC1000013	0.66
	SKNMC1000046	F-SKNMC1000046	0.72
15	SKNMC1000050	F-SKNMC1000050	0.47
	SKNMC1000091	F-SKNMC1000091	0.68
	THYR01000017	F-THYR01000017,	0.93
	THYR01000034	F-THYR01000034	0. 48
20	THYR01000040	F-THYR01000040	0. 32
	THYRO1000070	F-THYR01000070	0. 75
	THYR01000072	F-THYR01000072	0.94
	THYR01000107	F-THYR01000107	0.94
25	THYR01000121	F-THYR01000121	0.46
25	THYR01000124	F-THYR01000124	0. 37
	THYR01000163	F-THYR01000163	0. 45
	THYR01000173	F-THYR01000173	0. 94
	THYR01000197	F-THYR01000197	0. 78
30	THYR01000199	F-THYR01000199	0.55
	THYR01000206	F-THYR01000206	0. 59
	THYRO1000242	F-THYR01000242	0.69
	THYRO1000253	F-THYR01000253	0.48
35	THYR01000270	F-THYR01000270	0.85
	THYR01000288	F-THYR01000288	0.31
	THYR01000320	F-THYR01000320	0.94
	THYR01000327	F-THYR01000327	0.44
40	THYR01000343	F-THYR01000343	0.82
	THYR01000358	F-THYR01000358	0.89
	THYR01000368	F-THYR01000368	0.94
	THYR01000381	F-THYR01000381	0.64
45	THYR01000387	F-THYR01000387	0.57
	THYR01000394	F-THYR01000394	0.94
	THYR01000395	F-THYR01000395	0.91
	THYR01000401	F-THYR01000401 F-THYR01000488	0. 73 0. 94
50	THYR01000488 THYR01000501		0. 94
	THYR01000502	F-THYR01000501 F-THYR01000502	0.74
	THYR01000505	F-THYR01000505	0. 47
	THYR01000569	F-THYR01000569	0. 47
55	THYR01000570	F-THYR01000570	0.70
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		F TIP/004-00505	
		F-THYR01000585	0.60
	THYR01000605	F-THYR01000605	0. 79
	THYR01000637	F-THYR01000637	0.80
5	THYR01000662	F-THYR01000662	0. 77
	THYR01000666	F-THYR01000666	0. 47
	THYR01000676	F-THYR01000676	0. 94
	THYR01000684	F-THYR01000684	0. 72
10	THYR01000712	F-THYR01000712	0.94
	THYR01000715	F-THYR01000715	0.94
	THYR01000734	F-THYR01000734	0.44
	THYR01000756	F-THYR01000756	0. 94
15	THYR01000777	F-THYR01000777	0.73
	THYR01000783	F-THYR01000783	0.74
	THYR01000787	F-THYR01000787,	0. 67
	THYR01000829	F-THYR01000829	0. 43
	THYR01000855	F-THYR01000855	0. 91
20	THYR01000916	F-THYR01000916	0.64
	THYR01000926	F-THYR01000926	0.39
	THYR01000934	F-THYR01000934	0. 94
	THYR01000951	F-THYR01000951	0. 94
25	THYR01000952	F-THYR01000952	0. 67
	THYR01000983	F-THYR01000983	0.87
	THYR01000984	F-THYR01000984	0. 94
	THYR01000988	F-THYR01000988	0. 37
30	THYR01001033	F-THYR01001033	0.74
	THYR01001100	F-THYR01001100	0.74
	THYR01001120	F-THYR01001120	0.87
	THYR01001134	F-THYR01001134	0. 93
35	THYR01001142	F-THYR01001142	0. 61
	THYR01001173	F-THYR01001173	0.50
	THYR01001189	F-THYR01001189	0.40
	THYR01001204	F-THYR01001204	0. 94
	THYR01001213	F-THYR01001213	0.94
40	THYR01001271	F-THYR01001271	0.93
	THYR01001287	F-THYR01001287	0.32
	THYR01001313	F-THYR01001313	0. 53
	THYR01001320	F-THYR01001320	0. 68
45	THYR01001347	F-THYR01001347	0.68
	THYR01001363	F-THYR01001363	0.39
	THYR01001374	F-THYR01001374	0. 94
	THYR01001403	F-THYR01001403	0.49
50	THYR01001405	F-THYR01001405	0. 90
	THYR01001406	F-THYR01001406	0. 67
	THYR01001458		0. 94
	THYR01001487		0. 63
55	THYR01001537		0.94

	THYR01001584	F-THYR01001584	0.77
	THYR01001602	F-THYR01001602	0.64
	THYR01001605	F-THYR01001605	0.52
5	THYR01001637	F-THYR01001637_	0, 68
	THYR01001656	F-THYR01001656	0.82
	THYR01001661	F-THYR01001661	0.94
	THYR01001671	F-THYR01001671	0.94
10	THYR01001703	F-THYR01001703	0.87
	THYR01001721	F-THYR01001721	0.38
	THYR01001738	F-THYR01001738	0.51
	THYR01001793	F-THYR01001793	0.84
15	THYR01001809	F-THYR01001809	0.73
	THYR01001828	F-THYR01001828	0.88
	THYR01001854	F-THYR01001854,	0.47
	VESEN1000122	F-VESEN1000122	0.36
	Y79AA1000013	F-Y79AA1000013	0.94
20	Y79AA1000037	F-Y79AA1000037	0.94
	Y79AA1000059	F-Y79AA1000059	0.94
	Y79AA1000131	F-Y79AA1000131	0.56
	Y79AA1000181	F-Y79AA1000181	0.59
25	Y79AA1000202	F-Y79AA1000202	0, 45
	Y79AA1000214	F-Y79AA1000214	0.93
·	Y79AA1000230	F-Y79AA1000230	0.94
	Y79AA1000231	F-Y79AA1000231	0.36
30	Y79AA1000258	F-Y79AA1000258	0.36
	Y79AA1000268	F-Y79AA1000268	0.92
	Y79AA1000313	F-Y79AA1000313	0.91
	Y79AA1000328	F-Y79AA1000328	0.94
35	Y79AA1000342	F-Y79AA1000342	0.83
	Y79AA1000349	F-Y79AA1000349	0.50
	Y79AA1000355	F-Y79AA1000355	0.94
	Y79AA1000368	F-Y79AA1000368	0. 91
40	Y79AA1000420	F-Y79AA1000420	0. 94
	Y79AA1000469	F-Y79AA1000469	0.51
	Y79AA1000480	F-Y79AA1000480	0.94
	Y79AA1000539	F-Y79AA1000539	0. 74
	Y79AA1000540	F-Y79AA1000540	0.94
45	Y79AA1000560	F-Y79AA1000560	0. 93
	Y79AA1000574	F-Y79AA1000574	0. 87
	Y79AA1000589	F-Y79AA1000589	0.94
	Y79AA1000627	F-Y79AA1000627	0. 55
50	Y79AA1000705	F-Y79AA1000705	0.76
	Y79AA1000734	F-Y79AA1000734	0.60
	Y79AA1000748	F-Y79AA1000748	0. 45
	Y79AA1000752	F-Y79AA1000752	0.94
55	Y79AA1000774	F-Y79AA1000774	0.38

	Y79AA1000/82	F-Y/9AA1000782	0. 47
	Y79AA1000784	F-Y79AA1000784	0.93
_	Y79AA1000794	F-Y79AA1000794	0.67
5	Y79AA1000800	F-Y79AA10008Q0_	0.94
	Y79AA1000824	F-Y79AA1000824	0.90
	Y79AA1000827	F-Y79AA1000827	0. 94
	Y79AA1000833	F-Y79AA1000833	0.34
10	Y79AA1000850	F-Y79AA1000850	0. 92
	Y79AA1000962	F-Y79AA1000962	0. 69
	Y79AA1000966	F-Y79AA1000966	0.64
·	Y79AA1000968	F-Y79AA1000968	0. 64
15	Y79AA1000976	F-Y79AA1000976	0.66
	Y79AA1G00985	F-Y79AA1000985	0.66
	Y79AA1001023	F-Y79AA1001023	0. 33
	Y79AA1001041	F-Y79AA1001041	0. 90
20	Y79AA1001048	F-Y79AA1001048	0. 81
	Y79AA1001068	F-Y79AA1001068	0. 64
	Y79AA1001077	F-Y79AA1001077	0. 78
	Y79AA1001078	F-Y79AA1001078	0. 92
25	Y79AA1001105	F-Y79AA1001105	0.70
25	Y79AA1001145	F-Y79AA1001145	0.46
	Y79AA1001177	F-Y79AA1001177	0. 74
	Y79AA1001185	F-Y79AA1001185	0.74
	Y79AA1001211	F-Y79AA1001211	0. 94
30	Y79AA1001216	F-Y79AA1001216	0.61
	Y79AA1001228	F-Y79AA1001228	0. 56
	Y79AA1001233	F-Y79AA1001233	0. 83
	Y79AA1001236	F-Y79AA1001236	0. 32
35	Y79AA1001281	F-Y79AA1001281	0. 51
	Y79AA1001299	F-Y79AA1001299	0.94
	Y79AA1001312	F-Y79AA1001312	0. 53
	Y79AA1001323	F-Y79AA1001323	0. 76
40	Y79AA1001391	F-Y79AA1001391	0. 83
	Y79AA1001394	F-Y79AA1001394	0. 94
	Y79AA1001402	F~Y79AA1001402	0. 90
	Y79AA1001493	F-Y79AA1001493	0. 94
45	Y79AA1001511	F-Y79AA1001511	0. 47
	Y79AA1001533	F-Y79AA1001533	0. 92
	Y79AA1001541	F-Y79AA1001541	0. 32
	Y79AA1001548	F-Y79AA1001548	0. 50
50	Y79AA1001555	F-Y79AA1001555	0. 92
JU	Y79AA1001581	F-Y79AA1001581	0.94
	Y79AA1001585	F-Y79AA1001585	0.80
	Y79AA1001603	F-Y79AA1001603	0.94
	Y79AA1001613	F-Y79AA1001613	0.94
55	Y79AA1001647	F-Y79AA1001647	0.49

779AA1001679 F-Y79AA1001679 0, 94 779AA1001696 F-Y79AA1001696 0, 60 779AA1001705 F-Y79AA10017050, 67 779AA1001711 F-Y79AA1001711 0, 59 779AA1001805 F-Y79AA1001805 0, 79 779AA1001827 F-Y79AA1001827 0, 84 779AA1001827 F-Y79AA1001827 0, 84 779AA1001827 F-Y79AA1001827 0, 84 779AA1001827 F-Y79AA1001827 0, 84 779AA1001826 F-Y79AA1001827 0, 84 779AA1001826 F-Y79AA1001827 0, 84 779AA1001826 F-Y79AA1001827 0, 84 779AA1001827 F-Y79AA1001827 0, 82 779AA1001923 F-Y79AA1001827 0, 32 779AA1001923 F-Y79AA1001923 0, 32 779AA1002027 F-Y79AA100203 0, 65 779AA100203 F-Y79AA100203 0, 65 779AA100203 F-Y79AA100203 0, 89 779AA100203 F-Y79AA100203 0, 89 779AA100215 F-Y79AA100215 0, 31 779AA100215 F-Y79AA100215 0, 31 779AA1002204 F-Y79AA1002215 0, 76 779AA1002205 F-Y79AA1002204 0, 94 779AA1002206 F-Y79AA1002209 0, 91 779AA1002207 F-Y79AA1002209 0, 91 779AA1002208 F-Y79AA1002209 0, 91 779AA1002209 F-Y79AA1002209 0, 91 779AA1002209 F-Y79AA1002209 0, 91 779AA1002209 F-Y79AA1002209 0, 91 779AA1002209 F-Y79AA1002209 0, 91 779AA1002201 F-Y79AA1002209 0, 96 779AA1002209 F-Y79AA1002209 0, 96 779AA100231 F-Y79AA1002216 0, 65 779AA100231 F-Y79AA1002231 0, 85 779AA100231 F-Y79AA100231 0, 61 779AA1002416 F-Y79AA1002416 0, 77 779AA1002417 F-Y79AA1002416 0, 77 779AA1002417 F-Y79AA1002410 0, 77 779AA1002417 F-Y79AA1002410 0, 77 779AA1002417 F-Y79AA1002410 0, 77 779AA1002417 F-Y79AA1002407 0, 83 779AA1002418 F-Y79AA1002407 0, 83		Y79AA1001665	F-Y79AA1001665	0. 78
5         Y79AA1001705         F-Y79AA1001705         0. 67           Y79AA1001711         F-Y79AA1001711         0. 59           Y79AA1001805         F-Y79AA1001805         0. 79           Y79AA1001827         F-Y79AA1001827         0. 84           Y79AA1001846         F-Y79AA1001827         0. 84           Y79AA1001846         F-Y79AA1001866         0. 75           Y79AA1001827         F-Y79AA1001866         0. 75           Y79AA1001823         F-Y79AA1001823         0. 32           Y79AA1001963         F-Y79AA1001963         0. 51           Y79AA1002027         F-Y79AA1002023         0. 65           Y79AA1002033         F-Y79AA1002033         0. 65           Y79AA1002034         F-Y79AA1002039         0. 94           Y79AA1002035         F-Y79AA1002039         0. 99           Y79AA100215         F-Y79AA1002039         0. 99           Y79AA100215         F-Y79AA1002039         0. 99           Y79AA100215         F-Y79AA1002115         0. 32           Y79AA1002204         F-Y79AA1002209         0. 76           Y79AA1002205         F-Y79AA1002209         0. 91           Y79AA1002207         F-Y79AA1002209         0. 91           Y79AA1002209         F		Y79AA1001679		
5         Y79AA1001705         F-Y79AA1001751         0. 67           Y79AA1001711         F-Y79AA1001711         0. 59           Y79AA1001805         F-Y79AA1001805         0. 79           Y79AA1001827         F-Y79AA1001827         0. 84           Y79AA1001846         F-Y79AA1001866         0. 34           Y79AA1001866         F-Y79AA1001866         0. 75           Y79AA1001827         F-Y79AA1001866         0. 75           Y79AA1001866         F-Y79AA1001866         0. 59           Y79AA1001923         F-Y79AA1001923         0. 32           Y79AA1001923         F-Y79AA1001923         0. 51           Y79AA1002027         F-Y79AA1002083         0. 55           Y79AA100203         F-Y79AA1002083         0. 65           Y79AA100203         F-Y79AA1002093         0. 89           Y79AA100203         F-Y79AA1002093         0. 89           Y79AA1002115         F-Y79AA1002093         0. 90           Y79AA1002125         F-Y79AA1002115         0. 76           Y79AA1002204         F-Y79AA1002204         0. 94           Y79AA1002205         F-Y79AA1002209         0. 76           Y79AA1002207         F-Y79AA1002209         0. 76           Y79AA1002209         F		Y79AA1001696	F-Y79AA1001696	0.60
Y79AA1001711         F-Y79AA1001711         0. 59           Y79AA1001781         F-Y79AA1001781         0. 74           Y79AA1001805         F-Y79AA1001805         0. 79           Y79AA1001827         F-Y79AA1001827         0. 84           Y79AA1001846         F-Y79AA1001866         0. 75           Y79AA1001866         F-Y79AA1001866         0. 75           Y79AA1001923         F-Y79AA1001823         0. 32           Y79AA1001963         F-Y79AA1001963         0. 51           Y79AA1002027         F-Y79AA1002023         0. 65           Y79AA1002035         F-Y79AA1002033         0. 65           Y79AA1002037         F-Y79AA1002093         0. 94           Y79AA1002038         F-Y79AA1002093         F-Y79AA1002093           Y79AA1002155         F-Y79AA1002155         0. 76           Y79AA1002204         F-Y79AA1002204         0. 94           Y79AA1002205         F-Y79AA1002209         0. 91           Y79AA1002207         F-Y79AA1002209         0. 91           Y79AA1002208         F-Y79AA1002209         0. 76           Y79AA1002209         F-Y79AA1002209         0. 91           Y79AA1002209         F-Y79AA1002209         0. 91           Y79AA1002229         F-Y79AA1	5	Y79AA1001705	F-Y79AA1001705	0. 67
779AA1001805 F-Y79AA1001805 0.79 779AA1001827 F-Y79AA1001827 0.84 779AA1001846 F-Y79AA1001827 0.34 779AA1001866 F-Y79AA1001866 0.75 779AA1001874 F-Y79AA1001874 0.59 779AA1001923 F-Y79AA1001923 0.32 779AA1002027 F-Y79AA1001963 0.51 779AA1002027 F-Y79AA1002027, 0.39 779AA1002038 F-Y79AA1002027, 0.39 779AA1002038 F-Y79AA1002028, 0.65 779AA1002039 F-Y79AA1002030 0.89 779AA100215 F-Y79AA100215 0.76 779AA1002125 F-Y79AA1002139 0.32 779AA1002204 F-Y79AA1002139 0.32 779AA1002204 F-Y79AA100209 0.91 779AA1002204 F-Y79AA1002209 0.91 779AA1002209 F-Y79AA1002209 0.76 779AA1002219 F-Y79AA1002209 0.76 779AA1002229 F-Y79AA1002229 0.76 779AA1002234 F-Y79AA1002209 0.76 779AA1002235 F-Y79AA1002229 0.76 779AA1002236 F-Y79AA1002209 0.76 779AA1002237 F-Y79AA1002209 0.76 779AA1002238 F-Y79AA1002234 0.94 779AA1002361 F-Y79AA1002236 0.79 779AA1002361 F-Y79AA1002361 0.90 779AA1002361 F-Y79AA100239 0.94 779AA100246 F-Y79AA100239 0.94 779AA100246 F-Y79AA100239 0.94 779AA100247 F-Y79AA1002331 0.61 779AA1002431 F-Y79AA1002343 0.81 779AA1002442 F-Y79AA1002431 0.61 779AA1002442 F-Y79AA1002431 0.61 779AA1002442 F-Y79AA1002431 0.61 779AA1002442 F-Y79AA1002431 0.61 779AA1002442 F-Y79AA1002482 0.88 779AA1002482 F-Y79AA1002482 0.88		Y79AA1001711		
Y79AA1001827 F-Y79AA1001827 0. 84 Y79AA1001846 F-Y79AA1001846 0. 34 Y79AA1001866 F-Y79AA1001866 0. 75 Y79AA1001874 F-Y79AA1001874 0. 59 Y79AA1001923 F-Y79AA1001923 0. 32 Y79AA1002027 F-Y79AA1001963 0. 51 Y79AA1002027 F-Y79AA1002027, 0. 39 Y79AA1002088 F-Y79AA1002083 0. 65 Y79AA1002089 F-Y79AA1002083 0. 69 Y79AA1002093 F-Y79AA1002093 0. 89 Y79AA1002115 F-Y79AA1002115 0. 31 F-Y79AA1002125 F-Y79AA1002125 0. 76 Y79AA1002139 F-Y79AA1002139 0. 32 Y79AA1002204 F-Y79AA100204 0. 94 Y79AA1002208 F-Y79AA1002209 0. 91 Y79AA1002209 F-Y79AA1002209 0. 91 Y79AA1002219 F-Y79AA1002209 0. 60 Y79AA1002229 F-Y79AA1002229 0. 76 Y79AA1002246 F-Y79AA1002234 0. 94 Y79AA1002246 F-Y79AA1002234 0. 94 Y79AA1002258 F-Y79AA1002209 0. 76 Y79AA1002219 F-Y79AA1002209 0. 76 Y79AA1002210 F-Y79AA1002209 0. 91 Y79AA1002210 F-Y79AA1002210 0. 60 Y79AA1002210 F-Y79AA1002210 0. 60 Y79AA1002210 F-Y79AA1002210 0. 65 Y79AA1002210 F-Y79AA1002311 0. 92 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002407 F-Y79AA1002311 0. 92 Y79AA1002407 F-Y79AA1002311 0. 91 Y79AA1002407 F-Y79AA1002407 0. 83		Y79AA1001781	F-Y79AA1001781	0.74
Y79AA1001827 F-Y79AA1001827 0. 84 Y79AA1001846 F-Y79AA1001846 0. 34 Y79AA1001866 F-Y79AA1001866 0. 75 Y79AA1001874 F-Y79AA1001874 0. 59 Y79AA1001923 F-Y79AA1001923 0. 32 Y79AA1002027 F-Y79AA1001963 0. 51 Y79AA1002027 F-Y79AA1002027, 0. 39 Y79AA1002088 F-Y79AA1002083 0. 65 Y79AA1002089 F-Y79AA1002083 0. 69 Y79AA1002093 F-Y79AA1002093 0. 89 Y79AA1002115 F-Y79AA1002115 0. 31 F-Y79AA1002125 F-Y79AA1002125 0. 76 Y79AA1002139 F-Y79AA1002139 0. 32 Y79AA1002204 F-Y79AA100204 0. 94 Y79AA1002208 F-Y79AA1002209 0. 91 Y79AA1002209 F-Y79AA1002209 0. 91 Y79AA1002219 F-Y79AA1002209 0. 60 Y79AA1002229 F-Y79AA1002229 0. 76 Y79AA1002246 F-Y79AA1002234 0. 94 Y79AA1002246 F-Y79AA1002234 0. 94 Y79AA1002258 F-Y79AA1002209 0. 76 Y79AA1002219 F-Y79AA1002209 0. 76 Y79AA1002210 F-Y79AA1002209 0. 91 Y79AA1002210 F-Y79AA1002210 0. 60 Y79AA1002210 F-Y79AA1002210 0. 60 Y79AA1002210 F-Y79AA1002210 0. 65 Y79AA1002210 F-Y79AA1002311 0. 92 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002407 F-Y79AA1002311 0. 92 Y79AA1002407 F-Y79AA1002311 0. 91 Y79AA1002407 F-Y79AA1002407 0. 83	40	Y79AA1001805	F-Y79AA1001805	0. 79
15         Y79AA1001866         F-Y79AA1001874         0. 75           Y79AA1001874         F-Y79AA1001874         0. 59           Y79AA1001923         F-Y79AA1001923         0. 32           Y79AA1002027         F-Y79AA1002027, 0. 39           20         Y79AA1002083         F-Y79AA1002083         0. 65           Y79AA1002089         F-Y79AA1002093         0. 94           Y79AA1002093         F-Y79AA1002093         0. 89           Y79AA1002115         F-Y79AA1002115         0. 31           Y79AA1002125         F-Y79AA1002125         0. 76           Y79AA1002204         F-Y79AA1002204         0. 94           Y79AA1002203         F-Y79AA1002204         0. 94           Y79AA1002204         F-Y79AA1002204         0. 94           Y79AA1002203         F-Y79AA1002204         0. 94           Y79AA1002204         F-Y79AA1002204         0. 94           Y79AA1002205         F-Y79AA1002208         0. 76           Y79AA1002207         F-Y79AA1002209         0. 91           Y79AA1002209         F-Y79AA1002209         0. 76           Y79AA1002224         F-Y79AA1002224         0. 94           Y79AA1002234         F-Y79AA1002234         0. 94           Y79AA1002235	10	Y79AA1001827	F-Y79AA1001827	
15 Y79AA1001874 F-Y79AA1001874 0.59 Y79AA1001923 F-Y79AA1001923 0.32 Y79AA1001963 F-Y79AA1001963 0.51 Y79AA1002027 F-Y79AA1002027, 0.39 Y79AA1002083 F-Y79AA1002089 0.94 Y79AA1002093 F-Y79AA1002093 0.89 Y79AA1002115 F-Y79AA1002115 0.31 Y79AA100215 F-Y79AA1002125 0.76 Y79AA1002139 F-Y79AA1002139 0.32 Y79AA1002204 F-Y79AA1002139 0.32 Y79AA1002208 F-Y79AA1002208 0.76 Y79AA1002209 F-Y79AA1002209 0.91 Y79AA1002209 F-Y79AA1002209 0.91 Y79AA1002210 F-Y79AA1002210 0.60 Y79AA1002229 F-Y79AA1002229 0.76 Y79AA1002246 F-Y79AA1002234 0.94 Y79AA1002246 F-Y79AA1002234 0.94 Y79AA1002246 F-Y79AA1002238 0.79 Y79AA1002246 F-Y79AA1002238 0.79 Y79AA100234 F-Y79AA1002238 0.79 Y79AA100235 F-Y79AA1002236 0.65 Y79AA100236 F-Y79AA100237 0.52 Y79AA100236 F-Y79AA100231 0.92 Y79AA100237 F-Y79AA100231 0.92 Y79AA100239 F-Y79AA100231 0.92 Y79AA100246 F-Y79AA100231 0.92 Y79AA100247 F-Y79AA100240 0.83 Y79AA1002416 F-Y79AA100241 0.77 Y79AA1002416 F-Y79AA1002431 0.61 Y79AA1002431 F-Y79AA1002432 0.83 Y79AA1002431 F-Y79AA1002432 0.83 Y79AA1002431 F-Y79AA1002432 0.83 Y79AA1002431 F-Y79AA1002432 0.83 Y79AA1002433 F-Y79AA1002432 0.83 Y79AA1002442 F-Y79AA1002432 0.83 Y79AA1002442 F-Y79AA1002432 0.83 Y79AA1002433 F-Y79AA1002432 0.83		Y79AA1001846	F-Y79AA1001846	0.34
20       Y79AA1001923       F-Y79AA1001963       0. 51         20       Y79AA1002027       F-Y79AA1002027, 0. 39         20       Y79AA1002083       F-Y79AA1002083       0. 65         Y79AA1002089       F-Y79AA1002089       0. 94         Y79AA1002015       F-Y79AA1002093       0. 89         Y79AA100215       F-Y79AA1002115       0. 31         25       Y79AA1002125       F-Y79AA1002125       0. 76         Y79AA1002204       F-Y79AA1002204       0. 94         Y79AA1002204       F-Y79AA1002204       0. 94         Y79AA1002208       F-Y79AA1002204       0. 94         Y79AA1002209       F-Y79AA1002208       0. 76         Y79AA1002209       F-Y79AA1002209       0. 91         Y79AA1002209       F-Y79AA1002209       0. 91         Y79AA1002229       F-Y79AA1002220       0. 76         Y79AA1002234       F-Y79AA10022210       0. 60         Y79AA1002234       F-Y79AA1002234       0. 94         Y79AA1002234       F-Y79AA1002234       0. 94         Y79AA1002235       F-Y79AA1002236       0. 76         Y79AA1002236       F-Y79AA1002236       0. 70         Y79AA100237       F-Y79AA1002307       0. 52 <t< th=""><th></th><th>Y79AA1001866</th><th>F-Y79AA1001866</th><th>0. 75</th></t<>		Y79AA1001866	F-Y79AA1001866	0. 75
Y79AA1001963	15	Y79AA1001874	F-Y79AA1001874	0.59
20       Y79AA1002027       F-Y79AA1002083       0.65         Y79AA1002089       F-Y79AA1002089       0.94         Y79AA1002093       F-Y79AA1002093       0.89         Y79AA1002115       F-Y79AA1002115       0.31         25       Y79AA1002125       F-Y79AA1002125       0.76         Y79AA1002204       F-Y79AA1002204       0.94         Y79AA1002204       F-Y79AA1002204       0.94         Y79AA1002209       F-Y79AA1002209       0.91         Y79AA1002210       F-Y79AA1002209       0.91         Y79AA1002229       F-Y79AA1002229       0.76         Y79AA1002234       F-Y79AA1002234       0.94         Y79AA1002235       F-Y79AA1002234       0.94         Y79AA1002236       F-Y79AA1002236       0.55         Y79AA1002307       F-Y79AA1002230       0.52         Y79AA1002311       F-Y79AA1002311       0.92         Y79AA1002351       F-Y79AA1002361       0.90         Y79AA1002407       F-Y79AA1002		Y79AA1001923	F-Y79AA1001923	0.32
20 Y79AA1002083 F-Y79AA1002083 0.65 Y79AA1002089 F-Y79AA1002089 0.94 Y79AA100203 F-Y79AA1002033 0.89 Y79AA1002115 F-Y79AA1002115 0.31 Y79AA1002125 F-Y79AA1002125 0.76 Y79AA1002139 F-Y79AA1002139 0.32 Y79AA1002204 F-Y79AA1002204 0.94 Y79AA1002208 F-Y79AA1002204 0.94 Y79AA1002209 F-Y79AA1002209 0.91 Y79AA1002209 F-Y79AA1002209 0.76 Y79AA1002229 F-Y79AA1002229 0.76 Y79AA1002224 F-Y79AA1002224 0.94 Y79AA1002234 F-Y79AA1002234 0.94 Y79AA1002246 F-Y79AA1002234 0.94 Y79AA1002258 F-Y79AA1002258 0.79 Y79AA1002298 F-Y79AA1002258 0.79 Y79AA1002307 F-Y79AA1002258 0.83 Y79AA1002311 F-Y79AA1002311 0.92 Y79AA1002351 F-Y79AA1002351 0.85 Y79AA1002361 F-Y79AA1002361 0.90 Y79AA1002407 F-Y79AA1002361 0.90 Y79AA1002407 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002432 F-Y79AA1002433 0.81 Y79AA1002442 F-Y79AA1002432 0.88 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002482 F-Y79AA1002482 0.88		Y79AA1001963	F-Y79AA1001963	0.51
Y79AA1002089       F-Y79AA1002089       0.94         Y79AA1002093       F-Y79AA1002093       0.89         Y79AA1002115       F-Y79AA1002115       0.31         Y79AA1002125       F-Y79AA1002125       0.76         Y79AA1002204       F-Y79AA1002204       0.94         Y79AA1002208       F-Y79AA1002204       0.94         Y79AA1002209       F-Y79AA1002209       0.91         Y79AA1002210       F-Y79AA1002209       0.91         Y79AA1002229       F-Y79AA1002220       0.76         Y79AA1002229       F-Y79AA1002220       0.76         Y79AA1002229       F-Y79AA1002229       0.76         Y79AA1002234       F-Y79AA10022234       0.94         Y79AA1002234       F-Y79AA1002234       0.94         Y79AA1002235       F-Y79AA1002234       0.94         Y79AA1002238       F-Y79AA1002235       0.52         Y79AA1002307       F-Y79AA1002307       0.52         Y79AA1002311       F-Y79AA1002351       0.85         Y79AA1002361       F-Y79AA1002361       0.90         45       Y79AA1002407       F-Y79AA1002407       0.83         Y79AA1002407       F-Y79AA10024416       0.77         Y79AA1002431       F-Y79AA1002433       <		Y79AA1002027	F-Y79AA1002027	0.39
Y79AA1002093 F-Y79AA1002093 0.89 Y79AA1002115 F-Y79AA1002115 0.31 Y79AA1002125 F-Y79AA1002125 0.76 Y79AA1002139 F-Y79AA1002139 0.32 Y79AA1002204 F-Y79AA1002204 0.94 Y79AA1002208 F-Y79AA1002209 0.91 Y79AA1002209 F-Y79AA1002209 0.91 Y79AA1002210 F-Y79AA1002210 0.60 Y79AA1002229 F-Y79AA1002229 0.76 Y79AA1002234 F-Y79AA1002234 0.94 Y79AA1002246 F-Y79AA1002234 0.94 Y79AA1002258 F-Y79AA1002258 0.79 Y79AA1002258 F-Y79AA1002258 0.79 Y79AA1002307 F-Y79AA1002298 0.83 Y79AA1002311 F-Y79AA1002307 0.52 Y79AA1002311 F-Y79AA1002301 0.90 Y79AA1002307 F-Y79AA1002301 0.90 Y79AA1002307 F-Y79AA1002301 0.90 Y79AA1002407 F-Y79AA1002407 0.83	20	Y79AA1002083	F-Y79AA1002083	0.65
25       Y79AA1002115       F-Y79AA1002125       0. 76         Y79AA1002125       F-Y79AA1002139       0. 32         Y79AA1002204       F-Y79AA1002204       0. 94         Y79AA1002208       F-Y79AA1002208       0. 76         30       Y79AA1002209       F-Y79AA1002209       0. 91         Y79AA1002210       F-Y79AA1002210       0. 60         Y79AA1002229       F-Y79AA1002229       0. 76         35       Y79AA1002234       F-Y79AA1002234       0. 94         Y79AA1002246       F-Y79AA1002234       0. 94         Y79AA1002258       F-Y79AA1002258       0. 79         Y79AA1002307       F-Y79AA1002298       0. 83         40       Y79AA1002307       F-Y79AA1002307       0. 52         Y79AA1002311       F-Y79AA1002307       0. 52         Y79AA1002351       F-Y79AA1002351       0. 85         Y79AA1002361       F-Y79AA1002361       0. 90         45       Y79AA1002407       F-Y79AA1002407       0. 83         Y79AA1002416       F-Y79AA1002416       0. 77         Y79AA1002416       F-Y79AA1002431       0. 61         50       Y79AA1002433       F-Y79AA1002433       0. 81         Y79AA1002482       F-Y79AA1002482 <th></th> <th>Y79AA1002089</th> <th>F-Y79AA1002089</th> <th>0.94</th>		Y79AA1002089	F-Y79AA1002089	0.94
25       Y79AA1002125       F-Y79AA1002125       0.76         Y79AA1002139       F-Y79AA1002139       0.32         Y79AA1002204       F-Y79AA1002204       0.94         Y79AA1002208       F-Y79AA1002208       0.76         Y79AA1002209       F-Y79AA1002209       0.91         Y79AA1002210       F-Y79AA1002210       0.60         Y79AA1002229       F-Y79AA1002229       0.76         35       Y79AA1002234       F-Y79AA1002234       0.94         Y79AA1002246       F-Y79AA1002234       0.94         Y79AA1002246       F-Y79AA1002234       0.94         Y79AA1002298       F-Y79AA1002298       0.83         40       Y79AA1002307       F-Y79AA1002307       0.52         Y79AA1002311       F-Y79AA1002311       0.92         Y79AA1002351       F-Y79AA1002351       0.85         Y79AA1002361       F-Y79AA1002361       0.90         45       Y79AA1002407       F-Y79AA1002407       0.83         Y79AA1002416       F-Y79AA1002416       0.77         Y79AA1002431       F-Y79AA1002431       0.61         50       Y79AA1002433       F-Y79AA1002433       0.81         Y79AA1002482       F-Y79AA1002482       0.88		Y79AA1002093	F-Y79AA1002093	0.89
779AA1002125 F-779AA1002125 0.76 779AA1002139 F-Y79AA1002139 0.32 779AA1002204 F-Y79AA1002204 0.94 779AA1002208 F-Y79AA1002208 0.76 779AA1002209 F-Y79AA1002209 0.91 779AA1002210 F-Y79AA1002210 0.60 779AA1002229 F-Y79AA1002224 0.94 779AA1002234 F-Y79AA1002234 0.94 779AA1002246 F-Y79AA1002234 0.65 779AA1002258 F-Y79AA1002258 0.79 779AA1002298 F-Y79AA1002258 0.79 779AA1002307 F-Y79AA1002307 0.52 779AA1002311 F-Y79AA1002311 0.92 779AA1002351 F-Y79AA1002351 0.85 779AA1002361 F-Y79AA1002361 0.90 779AA1002407 F-Y79AA1002407 0.83 779AA1002416 F-Y79AA1002416 0.77 779AA1002416 F-Y79AA1002411 0.61 779AA1002433 F-Y79AA1002433 0.81 779AA1002472 F-Y79AA1002482 0.88 779AA1002482 F-Y79AA1002487 0.87		Y79AA1002115	F-Y79AA1002115	0. 31
779AA1002204 F-Y79AA1002204 0. 94 Y79AA1002208 F-Y79AA1002208 0. 76 Y79AA1002209 F-Y79AA1002209 0. 91 Y79AA1002210 F-Y79AA1002210 0. 60 Y79AA1002229 F-Y79AA1002229 0. 76 Y79AA1002234 F-Y79AA1002234 0. 94 Y79AA1002246 F-Y79AA1002234 0. 65 Y79AA1002258 F-Y79AA1002258 0. 79 Y79AA1002258 F-Y79AA1002258 0. 79 Y79AA1002307 F-Y79AA1002307 0. 52 Y79AA1002311 F-Y79AA1002307 0. 52 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002351 F-Y79AA1002351 0. 85 Y79AA1002361 F-Y79AA1002361 0. 90  45 Y79AA1002361 F-Y79AA1002361 0. 90 45 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002416 F-Y79AA1002416 0. 77 Y79AA1002417 F-Y79AA1002431 0. 61 Y79AA1002431 F-Y79AA1002431 0. 61 Y79AA1002432 F-Y79AA1002432 0. 88 Y79AA1002482 F-Y79AA1002482 0. 88 Y79AA1002487 F-Y79AA1002487 0. 87	25	Y79AA1002125	F-Y79AA1002125	0. 76
779AA1002208 F-Y79AA1002208 0. 76 779AA1002209 F-Y79AA1002209 0. 91 779AA1002210 F-Y79AA1002210 0. 60 779AA1002229 F-Y79AA1002229 0. 76 779AA1002234 F-Y79AA1002234 0. 94 779AA1002246 F-Y79AA1002246 0. 65 779AA1002258 F-Y79AA1002258 0. 79 779AA1002298 F-Y79AA1002298 0. 83 779AA1002307 F-Y79AA1002307 0. 52 779AA1002311 F-Y79AA1002311 0. 92 779AA1002351 F-Y79AA1002351 0. 85 779AA1002361 F-Y79AA1002361 0. 90 779AA1002361 F-Y79AA1002361 0. 90 779AA1002407 F-Y79AA1002407 0. 83		Y79AA1002139	F-Y79AA1002139	0. 32
30       Y79AA1002209       F-Y79AA1002209       0. 91         Y79AA1002210       F-Y79AA1002210       0. 60         Y79AA1002229       F-Y79AA1002229       0. 76         Y79AA1002234       F-Y79AA1002234       0. 94         Y79AA1002246       F-Y79AA1002246       0. 65         Y79AA1002258       F-Y79AA1002258       0. 79         Y79AA1002298       F-Y79AA1002298       0. 83         40       Y79AA1002307       F-Y79AA1002307       0. 52         Y79AA1002311       F-Y79AA1002311       0. 92         Y79AA1002351       F-Y79AA1002351       0. 85         Y79AA1002361       F-Y79AA1002361       0. 90         45       Y79AA1002399       F-Y79AA1002399       0. 94         Y79AA1002407       F-Y79AA1002407       0. 83         Y79AA1002416       F-Y79AA1002431       0. 61         50       Y79AA1002433       F-Y79AA1002433       0. 81         Y79AA1002472       F-Y79AA1002482       0. 88         Y79AA1002482       F-Y79AA1002482       0. 88         Y79AA1002487       F-Y79AA1002487       0. 87		Y79AA1002204	F-Y79AA1002204	0.94
779AA1002209 F-Y79AA1002209 0. 91 Y79AA1002210 F-Y79AA1002210 0. 60 Y79AA1002229 F-Y79AA1002229 0. 76 Y79AA1002234 F-Y79AA1002234 0. 94 Y79AA1002246 F-Y79AA1002246 0. 65 Y79AA1002258 F-Y79AA1002258 0. 79 Y79AA1002298 F-Y79AA1002298 0. 83 Y79AA1002307 F-Y79AA1002307 0. 52 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002351 F-Y79AA1002351 0. 85 Y79AA1002361 F-Y79AA1002361 0. 90 Y79AA1002407 F-Y79AA1002399 0. 94 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002407 F-Y79AA1002416 0. 77 Y79AA1002416 F-Y79AA1002431 0. 61 Y79AA1002431 F-Y79AA1002431 0. 61 Y79AA1002432 F-Y79AA1002432 0. 39 Y79AA1002482 F-Y79AA1002482 0. 88 Y79AA1002487 F-Y79AA1002487 0. 87	30	Y79AA1002208	F-Y79AA1002208	0. 76
779AA1002229 F-Y79AA1002229 0. 76 779AA1002234 F-Y79AA1002234 0. 94 779AA1002246 F-Y79AA1002246 0. 65 779AA1002258 F-Y79AA1002258 0. 79 779AA1002298 F-Y79AA1002298 0. 83 779AA1002307 F-Y79AA1002307 0. 52 779AA1002311 F-Y79AA1002311 0. 92 779AA1002351 F-Y79AA1002351 0. 85 779AA1002361 F-Y79AA1002361 0. 90 779AA1002399 F-Y79AA1002399 0. 94 779AA1002407 F-Y79AA1002407 0. 83 779AA1002416 F-Y79AA1002416 0. 77 779AA1002431 F-Y79AA1002431 0. 61 779AA1002432 F-Y79AA1002432 0. 88 779AA1002482 F-Y79AA1002482 0. 88 779AA1002487 F-Y79AA1002487 0. 87	30		F-Y79AA1002209	
779AA1002234 F-Y79AA1002234 0. 94 Y79AA1002246 F-Y79AA1002246 0. 65 Y79AA1002258 F-Y79AA1002258 0. 79 Y79AA1002298 F-Y79AA1002298 0. 83 Y79AA1002307 F-Y79AA1002307 0. 52 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002351 F-Y79AA1002351 0. 85 Y79AA1002361 F-Y79AA1002361 0. 90 Y79AA1002399 F-Y79AA1002399 0. 94 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002416 F-Y79AA1002416 0. 77 Y79AA1002431 F-Y79AA1002431 0. 61 Y79AA1002432 F-Y79AA1002432 0. 88 Y79AA1002482 F-Y79AA1002487 0. 87			F-Y79AA1002210	
Y79AA1002246 F-Y79AA1002246 0. 65 Y79AA1002258 F-Y79AA1002258 0. 79 Y79AA1002298 F-Y79AA1002298 0. 83 Y79AA1002307 F-Y79AA1002307 0. 52 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002351 F-Y79AA1002351 0. 85 Y79AA1002361 F-Y79AA1002361 0. 90 Y79AA1002399 F-Y79AA1002399 0. 94 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002416 F-Y79AA1002416 0. 77 Y79AA1002431 F-Y79AA1002431 0. 61 Y79AA1002433 F-Y79AA1002433 0. 81 Y79AA1002472 F-Y79AA1002472 0. 39 Y79AA1002482 F-Y79AA1002487 0. 87				
Y79AA1002258 F-Y79AA1002258 0. 79 Y79AA1002298 F-Y79AA1002298 0. 83 Y79AA1002307 F-Y79AA1002307 0. 52 Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002351 F-Y79AA1002351 0. 85 Y79AA1002361 F-Y79AA1002361 0. 90 Y79AA1002399 F-Y79AA1002399 0. 94 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002416 F-Y79AA1002416 0. 77 Y79AA1002431 F-Y79AA1002431 0. 61 Y79AA1002432 F-Y79AA1002432 0. 88 Y79AA1002482 F-Y79AA1002482 0. 88 Y79AA1002487 F-Y79AA1002487 0. 87	35			
Y79AA1002298 F-Y79AA1002298 0.83 Y79AA1002307 F-Y79AA1002307 0.52 Y79AA1002311 F-Y79AA1002311 0.92 Y79AA1002351 F-Y79AA1002351 0.85 Y79AA1002361 F-Y79AA1002361 0.90 Y79AA1002399 F-Y79AA1002399 0.94 Y79AA1002407 F-Y79AA1002407 0.83 Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002432 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002487 0.87				
Y79AA1002307 F-Y79AA1002307 0.52 Y79AA1002311 F-Y79AA1002311 0.92 Y79AA1002351 F-Y79AA1002351 0.85 Y79AA1002361 F-Y79AA1002361 0.90 Y79AA1002399 F-Y79AA1002399 0.94 Y79AA1002407 F-Y79AA1002407 0.83 Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002487 0.87				
Y79AA1002311 F-Y79AA1002311 0. 92 Y79AA1002351 F-Y79AA1002351 0. 85 Y79AA1002361 F-Y79AA1002361 0. 90 Y79AA1002399 F-Y79AA1002399 0. 94 Y79AA1002407 F-Y79AA1002407 0. 83 Y79AA1002416 F-Y79AA1002416 0. 77 Y79AA1002431 F-Y79AA1002431 0. 61 Y79AA1002433 F-Y79AA1002433 0. 81 Y79AA1002472 F-Y79AA1002472 0. 39 Y79AA1002482 F-Y79AA1002482 0. 88 Y79AA1002487 F-Y79AA1002487 0. 87				
Y79AA1002351 F-Y79AA1002351 0.85 Y79AA1002361 F-Y79AA1002361 0.90 Y79AA1002399 F-Y79AA1002399 0.94 Y79AA1002407 F-Y79AA1002407 0.83 Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002487 0.87	40			
Y79AA1002361 F-Y79AA1002361 0.90 Y79AA1002399 F-Y79AA1002399 0.94 Y79AA1002407 F-Y79AA1002407 0.83 Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87	·			
Y79AA1002399 F-Y79AA1002399 0.94 Y79AA1002407 F-Y79AA1002407 0.83 Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87				
Y79AA1002407 F-Y79AA1002407 0.83 Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87				
Y79AA1002416 F-Y79AA1002416 0.77 Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87	45			
Y79AA1002431 F-Y79AA1002431 0.61 Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87				
Y79AA1002433 F-Y79AA1002433 0.81 Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87				
Y79AA1002472 F-Y79AA1002472 0.39 Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87	50			
Y79AA1002482 F-Y79AA1002482 0.88 Y79AA1002487 F-Y79AA1002487 0.87	-			
Y79AA1002487 F-Y79AA1002487 0.87				
	55		1 1/3MA100240/	U. 01

### Table 10

Selected clones having the maximal ATGpr1 score of not more than 0.3 (1857 clones including the 9 clones whose 5'-end sequence does not contain the ATG codon. Blank of the maximal ATGpr1 score indicates that clones whose 5'-end sequence does not contain the ATG codon).

	clone	name of	maximal
	name	sequence	ATGprl
5		-	score
	HEMBA100004	2 F-HEMBA1000	0.08
	HEMBA100011	1 F-HEMBA1000	0111 0.1
10	HEMBA100018	O F-HEMBA1000	0180 0.09
	HEMBA100021	3 F-HEMBA1000	0213 0.21
	HEMBA100024	3 F-HEMBA1000	0243 0.1
•	HEMBA100025	1 F-HEMBA1000	0251 0.08
15	HEMBA100026	4 F-HEMBA1000	0264 0.08
	HEMBA100028	O F-HEMBA1000	0280 0.13
	HEMBA100028	2 F-HEMBA1000	0282 , 0. 1
	HEMBA100033	3 F-HEMBA1000	0333 0.17
20	HEMBA100033	8 F-HEMBA1000	0338 0.16
20	HEMBA100035	1 F-HEMBA1000	0351 0.17
	HEMBA100035	55 F-HEMBA1000	0355 0.11
	HEMBA100035	7 F-HEMBA1000	0357 0.08
0.5	HEMBA100036	66 F-HEMBA100	0366 0.11
25	HEMBA100037	6 F-HEMBA100	0376 0.1
	HEMBA100039	00 F-HEMBA100	0390 0.11
	HEMBA100041	11 F-HEMBA1000	0411 0.07
	HEMBA100041	18 F-HEMBA100	0418 0.06
30	HEMBA100042	22 F-HEMBA100	0422 0.07
	HEMBA100042	28 F-HEMBA1000	0428 0.08
	HEMBA100043	34 F-HEMBA1000	0434 0.11
	HEMBA100044	12 F-HEMBA100	0442 0.1
35	HEMBA100045	59 F-HEMBA1000	
	HEMBA100046	64 F-HEMBA1000	
	HEMBA100046	69 F-HEMBA100	0469 0.07
	HEMBA100050	04 F-HEMBA100	
40	HEMBA100051		
	HEMBA100051		
	HEMBA100054	•	
	HEMBA100054		
45	HEMBA100055		
	HEMBA100056		
	HEMBA100056		
	HEMBA100057		
50	HEMBA100060		
	HEMBA100062		
	HEMBA100065		
	HEMBA100066		
55	HEMBA100067	•	
	HEMBA100070	02 F-HEMBA100	0702 0.12

	HEMBA1000705	F-HEMBA1000705	0.06
	HEMBA1000722	F-HEMBA1000722	0. 17
	HEMBA1000726	F-HEMBA1000726	0.08
5	HEMBA1000747	F-HEMBA1000747	
	HEMBA1000749	F-HEMBA1000749	0.09
	HEMBA1000769	F-HEMBA1000769	0. 19
	HEMBA1000773	F-HEMBA1000773	0.08
10	HEMBA1000774	F-HEMBA1000774	0. 17
	HEMBA1000791	F-HEMBA1000791	0, 08
	HEMBA1000822	F-HEMBA1000822	0. 19
	HEMBA1000827	F-HEMBA1000827	0. 09
15	HEMBA1000843	F-HEMBA1000843	0. 12
15	HEMBA1000852	F-HEMBA1000852	0. 16
	HEMBA1000870	F-HEMBA1000870	
	HEMBA1000876	F-HEMBA1000876	0. 15
	HEMBA1000908	F-HEMBA1000908	0. 17
20	HEMBA1000934	F-HEMBA1000934	0. 2
	HEMBA1000943	F-HEMBA1000943	0.08
	HEMBA1000960	F-HEMBA1000960	0.09
	HEMBA1000972	F-HEMBA1000972	0.12
25	HEMBA1000974	F-HEMBA1000974	0.08
	HEMBA1000985	F-HEMBA1000985	0.06
	HEMBA1000991	F-HEMBA1000991	0. 23
	HEMBA1001007	F-HEMBA1001007	0.06
30	HEMBA1001017	F-HEMBA1001017	0.06
	HEMBA1001019	F-HEMBA1001019	0.08
	HEMBA1001020	F-HEMBA1001020	0.15
	HEMBA1001024	F-HEMBA1001024	0.1
35	HEMBA1001026	F-HEMBA1001026	0. 13
	HEMBA1001051	F-HEMBA1001051	0. 27
	HEMBA1001060	F-HEMBA1001060	0.11
	HEMBA1001071	F-HEMBA1001071	0.1
40	HEMBA1001077	F-HEMBA1001077	0. 07
	HEMBA1001094	F-HEMBA1001094	0. 07
	HEMBA1001099	F-HEMBA1001099	0.08
	HEMBA1001121	F-HEMBA1001121	0. 25
45	HEMBA1001123	F-HEMBA1001123	0. 05
	HEMBA1001172	F-HEMBA1001172	0. 08
	HEMBA1001208	F-HEMBA1001208	0. 11
	HEMBA1001226	F-HEMBA1001226	0. 27
50	HEMBA1001265	F-HEMBA1001265	0.06
	HEMBA1001294	F-HEMBA1001294	0.12
	HEMBA1001299	F-HEMBA1001299	0. 2
	HEMBA1001319	F-HEMBA1001319	0. 3
55 ·	HEMBA1001323		0. 07
	HEMBA1001327	F-HEMBA1001327	0.08

	HEMBA1001330	F-HEMBA1001330	0.11
	HEMBA1001375	F-HEMBA1001375	0. 15
	HEMBA1001383	F-HEMBA1001383	0.11
5	HEMBA1001391	F-HEMBA1001391_	0, 21
	HEMBA1001411	F-HEMBA1001411	0.06
	HEMBA1001432	F-HEMBA1001432	0.07
	HEMBA1001433	F-HEMBA1001433	0. 24
10	HEMBA1001435	F-HEMBA1001435	0.16
	HEMBA1001442	F-HEMBA1001442	0. 22
	HEMBA1001454	F-HEMBA1001454	0.11
	HEMBA1001463	F-HEMBA1001463	0.08
15	HEMBA1001478	F-HEMBA1001478	0.13
	HEMBA1001515	F-HEMBA1001515	0.3
	HEMBA1001517	F-HEMBA1001517	0.07
	HEMBA1001522	F-HEMBA1001522	0.3
20	HEMBA1001557	F-HEMBA1001557	0.06
	HEMBA1001566	F-HEMBA1001566	0.1
	HEMBA1001585	F-HEMBA1001585	0.09
	HEMBA1001589	F-HEMBA1001589	0.06
25	HEMBA1001608	F-HEMBA1001608	0.17
	HEMBA1001636	F-HEMBA1001636	0.15
	HEMBA1001651	F-HEMBA1001651	0. 23
	HEMBA1001658	F-HEMBA1001658	0.17
30	HEMBA1001675	F-HEMBA1001675	0.11
	HEMBA1001678	F-HEMBA1001678	0.06
	HEMBA1001681 HEMBA1001709	F-HEMBA1001681 F-HEMBA1001709	0. 1 0. 29
	HEMBA1001712	F-HEMBA1001712	0. 29
35	HEMBA1001718	F-HEMBA1001718	0. 07
•	HEMBA1001734	F-HEMBA1001734	0. 07
	HEMBA1001745	F-HEMBA1001745	0. 09
	HEMBA1001761	F-HEMBA1001761	0. 09
40	HEMBA1001784	F-HEMBA1001784	0. 11
	HEMBA1001791	F-HEMBA1001791	0.15
	HEMBA1001803	F-HEMBA1001803	0. 21
	HEMBA1001808	F-HEMBA1001808	0.18
45	HEMBA1001820	F-HEMBA1001820	0.08
40	HEMBA1001835	F-HEMBA1001835	0. 29
	HEMBA1001844	F-HEMBA1001844	0.11
	HEMBA1001861	F-HEMBA1001861	0. 2
50	HEMBA1001888	F-HEMBA1001888	0.18
50	HEMBA1001918	F-HEMBA1001918	0.09
	HEMBA1001940	F-HEMBA1001940	0. 21
	HEMBA1001942	F-HEMBA1001942	0.17
	HEMBA1001945	F-HEMBA1001945	0.06
55 .	HEMBA1001960	F-HEMBA1001960	

	HEMBA1001962	F-HEMBA1001962	0. 08
	HEMBA1001964	F-HEMBA1001964	0.15
	HEMBA1001979	F-HEMBA1001979	0. 07
5	HEMBA1001991	F-HEMBA1001991	0.08
	HEMBA1002003	F-HEMBA1002003	0, 11
	HEMBA1002008	F-HEMBA1002008	0.08
	HEMBA1002022	F-HEMBA1002022	0.08
10	HEMBA1002039	F-HEMBA1002039	0. 2
10	HEMBA1002100	F-HEMBA1002100	0.07
	HEMBA1002113	F-HEMBA1002113	0.09
	HEMBA1002139	F-HEMBA1002139	0. 2
15	HEMBA1002144	F-HEMBA1002144	0.11
15	HEMBA1002153	F-HEMBA1002153	0.09
	HEMBA1002160	F-HEMBA1002160,	0. 15
	HEMBA1002162	F-HEMBA1002162	0.19
	HEMBA1002166	F-HEMBA1002166	0.06
20	HEMBA1002185	F-HEMBA1002185	0.11
	HEMBA1002204	F-HEMBA1002204	0.09
	HEMBA1002226	F-HEMBA1002226	0.13
	HEMBA1002253	F-HEMBA1002253	0.05
25	HEMBA1002257	F-HEMBA1002257	0.09
	HEMBA1002270	F-HEMBA1002270	0.06
	HEMBA1002321	F-HEMBA1002321	0. 07
	HEMBA1002328	F-HEMBA1002328	0.06
30	HEMBA1002337	F-HEMBA1002337	0. 11
	HEMBA1002348	F-HEMBA1002348	0. 12
	HEMBA1002349	F-HEMBA1002349	0.05
	HEMBA1002381	F-HEMBA1002381	0.09
35	HEMBA1002389	F-HEMBA1002389	0. 14
	HEMBA1002486	F-HEMBA1002486	0. 1
	HEMBA1002498	F-HEMBA1002498	0. 17
	HEMBA1002538	F-HEMBA1002538	0. 27
40	HEMBA1002552	F-HEMBA1002552	0. 23
•	HEMBA1002555	F-HEMBA1002555	0. 11
	HEMBA1002558	F-HEMBA1002558	0.11
	HEMBA1002561	F-HEMBA1002561	0.11
45	HEMBA1002590	F-HEMBA1002590	0.06
	HEMBA1002592	F-HEMBA1002592	0. 17
	HEMBA1002621	F-HEMBA1002621	0. 12
·	HEMBA1002628	F-HEMBA1002628	0.06
50	HEMBA1002629	F-HEMBA1002629	0. 23
•	HEMBA1002645	F-HEMBA1002645	0. 12
	HEMBA1002651	F-HEMBA1002651	0.08
	HEMBA1002659	F-HEMBA1002659	0. 28
55	HEMBA1002661 HEMBA1002666	F-HEMBA1002661 F-HEMBA1002666	0. 28
55	TILMDATOUZOOD	TIEMPATOVZ000	0. 16

	HEMBA1002678	F-HEMBA1002678	0 07
	HEMBA1002679	F-HEMBA1002679	0. 07
	HEMBA1002712		0. 14
5		F-HEMBA1002712	0. 26
·	HEMBA1002716	F-HEMBA1002716	0.14
		F-HEMBA1002728	0.09
	HEMBA1002730	F-HEMBA1002730	0.07
	HEMBA1002742	F-HEMBA1002742	0.09
10	HEMBA1002748	F-HEMBA1002748	0.17
	HEMBA1002780	F-HEMBA1002780	0.1
	HEMBA1002801	F-HEMBA1002801	0.1
	HEMBA1002826	F-HEMBA1002826	0.1
15	HEMBA1002833	F-HEMBA1002833	0. 14
	HEMBA1002886	F-HEMBA1002886	0.09
	HEMBA1002896	F-HEMBA1002896,	0.05
	HEMBA1002921	F-HEMBA1002921	0. 25
20	HEMBA1002924	F-HEMBA1002924	0.12
	HEMBA1002934	F-HEMBA1002934	0. 2
	HEMBA1002944	F-HEMBA1002944	0.12
	HEMBA1002968	F-HEMBA1002968	0.12
25	HEMBA1003034	F-HEMBA1003034	0.06
20	HEMBA1003037	F-HEMBA1003037	0. 11
	HEMBA1003064	F-HEMBA1003064	0.09
	HEMBA1003071	F-HEMBA1003071	0. 27
	HEMBA1003083	F-HEMBA1003083	0.18
30	HEMBA1003086	F-HEMBA1003086	0.06
	HEMBA1003098	F-HEMBA1003098	0.09
	HEMBA1003133	F-HEMBA1003133	0. 12
	HEMBA1003142	F-HEMBA1003142	0. 29
35	HEMBA1003166	F-HEMBA1003166	0. 1
	HEMBA1003197	F-HEMBA1003197	0. 1
	HEMBA1003202	F-HEMBA1003202	0. 17
	HEMBA1003204	F-HEMBA1003204	0. 13
40	HEMBA1003212	F-HEMBA1003212	0. 13
	HEMBA1003220	F-HEMBA1003220	0. 14
	HEMBA1003229	F-HEMBA1003229	0.19
	HEMBA1003273	F-HEMBA1003273	0. 1
45	HEMBA1003276	F-HEMBA1003276	0.12
•	HEMBA1003278	F-HEMBA1003278	0.09
	HEMBA1003296	F-HEMBA1003296	0.07
	HEMBA1003304	F-HEMBA1003304	0.09
50	HEMBA1003309	F-HEMBA1003309	0.07
50	HEMBA1003314	F-HEMBA1003314	0. 21
	HEMBA1003328	F-HEMBA1003328	0.13
	HEMBA1003330	F-HEMBA1003330	0.08
	HEMBA1003348	F-HEMBA1003348	0.17
55	HEMBA1003373	F-HEMBA1003373	0. 21

}	IEMBA1003376	F-HEMBA1003376	0.14
ł	IEMBA1003384	F-HEMBA1003384	0.07
ł	IEMBA1003531	F-HEMBA1003531	0.2
5	EMBA1003548	F-HEMBA1003548	0.08
1	EMBA1003571	F-HEMBA1003571	0, 13
1	EMBA1003579	F-HEMBA1003579	0.13
	HEMBA1003595	F-HEMBA1003595	0.08
	HEMBA1003597	F-HEMBA1003597	0.1
	HEMBA1003598	F-HEMBA1003598	0.12
	HEMBA1003622	F-HEMBA1003622	0.08
	HEMBA1003630	F-HEMBA1003630	0.08
15	HEMBA1003637	F-HEMBA1003637	0.07
13	HEMBA1003640	F-HEMBA1003640	0.16
	HEMBA1003656	F-HEMBA1003656,	0.13
	HEMBA1003714	F-HEMBA1003714	0.12
	HEMBA1003715	F-HEMBA1003715	0.1
20	HEMBA1003725	F-HEMBA1003725	0.08
	HEMBA1003733	F-HEMBA1003733	0. 2
	HEMBA1003758	F-HEMBA1003758	0.06
	HEMBA1003773	F-HEMBA1003773	0.11
25	HEMBA1003784	F-HEMBA1003784	0.09
	HEMBA1003856	F-HEMBA1003856	0. 25
	HEMBA1003885	F-HEMBA1003885	0.09
	HEMBA1003902	F-HEMBA1003902	0.08
30	HEMBA1003908	F-HEMBA1003908	0.08
	HEMBA1003926	F-HEMBA1003926	0.09
	HEMBA1003937	F-HEMBA1003937	0. 2
	HEMBA1003939	F-HEMBA1003939	0.12
35	HEMBA1003942	F-HEMBA1003942	0. 1
	HEMBA1003950	F-HEMBA1003950	0.09
	HEMBA1003958	F-HEMBA1003958	0. 07
	HEMBA1003976	F-HEMBA1003976	0. 28
40	HEMBA1003987	F-HEMBA1003987	0. 24
	HEMBA1004000	F-HEMBA1004000	0.16
	HEMBA1004012	F-HEMBA1004012	0.12
	HEMBA1004015	F-HEMBA1004015	0. 23
45	HEMBA1004024	F-HEMBA1004024	0.13
	HEMBA1004038	F-HEMBA1004038	0.09
	HEMBA1004042	F-HEMBA1004042	0.09
	HEMBA1004045	F-HEMBA1004045	0.09
50	HEMBA1004049	F-HEMBA1004049	0.09
30	HEMBA1004132	F-HEMBA1004132	0.08
	HEMBA1004138	F-HEMBA1004138	0.07
	HEMBA1004164	•	0.08
	HEMBA1004225		0.1
55	HEMBA1004241	F-HEMBA1004241	0.08

	HEMBA1004267	F-HEMBA1004267	0.09
	HEMBA1004272	F-HEMBA1004272	0.09
	HEMBA1004295	F-HEMBA1004295	0.19
5	HEMBA1004306	F-HEMBA1004306	0.07
	HEMBA1004312	F-HEMBA1004312	0. 26
	HEMBA1004323	F-HEMBA1004323	0.13
	HEMBA1004334	F-HEMBA1004334	0.06
10	HEMBA1004354	F-HEMBA1004354	0. 1
	HEMBA1004356	F-HEMBA1004356	0. 07
	HEMBA1004366	F-HEMBA1004366	0.14
	HEMBA1004394	F-HEMBA1004394	0.08
15	HEMBA1004396	F-HEMBA1004396	0.05
	HEMBA1004405	F-HEMBA1004405	0. 17
	HEMBA1004429	F-HEMBA1004429,	
	HEMBA1004433	F-HEMBA1004433	0. 29
	HEMBA1004460	F-HEMBA1004460	0. 11
20	HEMBA1004461	F-HEMBA1004461	0.11
	HEMBA1004482	F-HEMBA1004482	0, 15
	HEMBA1004506	F-HEMBA1004506	0.05
	HEMBA1004538	F-HEMBA1004538	0.15
25	HEMBA1004577	F-HEMBA1004577	0. 17
	HEMBA1004586	F-HEMBA1004586	0.06
	HEMBA1004617	F-HEMBA1004617	0.11
	HEMBA1004629	F-HEMBA1004629	0.11
30	HEMBA1004631	F-HEMBA1004631	0.1
	HEMBA1004666	F-HEMBA1004666	0.08
	HEMBA1004670	F-HEMBA1004670	0.15
	HEMBA1004672	F-HEMBA1004672	0.14
35	HEMBA1004730	F-HEMBA1004730	0. 15
	HEMBA1004733	F-HEMBA1004733	0, 12
	HEMBA1004748	F-HEMBA1004748	0.1
	HEMBA1004770	F-HEMBA1004770	0.15
40	HEMBA1004778	F-HEMBA1004778	0.14
	HEMBA1004803	F-HEMBA1004803	0.13
	HEMBA1004807	F-HEMBA1004807	0.18
	HEMBA1004816	F-HEMBA1004816	0.06
45	HEMBA1004820	F-HEMBA1004820	0.11
45	HEMBA1004865	F-HEMBA1004865	0.17
	HEMBA1004880	F-HEMBA1004880	0. 28
	HEMBA1004900	F-HEMBA1004900	0.11
	HEMBA1004909	F-HEMBA1004909	0.11
50	HEMBA1004918	F-HEMBA1004918	0.09
	HEMBA1004956	F-HEMBA1004956	0.05
	HEMBA1004960	F-HEMBA1004960	0. 21
	HEMBA1004978	F-HEMBA1004978	0.1
55	HEMBA1004983	F-HEMBA1004983	0. 28

	HEMBA1004995	F~HEMBA1004995	0. 24
	HEMBA1005008	F-HEMBA1005008	0. 07
	HEMBA1005039	F-HEMBA1005039	0.08
5	HEMBA1005062	F-HEMBA1005062	0.11
	HEMBA1005123	F-HEMBA1005123	0. 07
	HEMBA1005152	F-HEMBA1005152	0.11
	HEMBA1005159	F-HEMBA1005159	0.1
10	HEMBA1005232	F-HEMBA1005232	0.1
	HEMBA1005241	F-HEMBA1005241	0.14
	HEMBA1005251	F-HEMBA1005251	0. 29
	HEMBA1005274	F-HEMBA1005274	0.06
15	HEMBA1005275	F-HEMBA1005275	0. 12
	HEMBA1005293	F-HEMBA1005293	0.06
	HEMBA1005304	F-HEMBA1005304,	
	HEMBA1005311	F-HEMBA1005311	0. 2
20	HEMBA1005315	F-HEMBA1005315	0.07
	HEMBA1005318	F-HEMBA1005318	0.08
	HEMBA1005353	F-HEMBA1005353	0.12
	HEMBA1005359	F-HEMBA1005359	0. 15
	HEMBA1005372	F-HEMBA1005372	0. 07
25	HEMBA1005374	F-HEMBA1005374	0.09
	HEMBA1005389	F-HEMBA1005389	0.1
	HEMBA1005408	F-HEMBA1005408	0.09
	HEMBA1005410	F-HEMBA1005410	0.07
30	HEMBA1005426	F-HEMBA1005426	0.16
	HEMBA1005443	F-HEMBA1005443	0. 17
	HEMBA1005447	F-HEMBA1005447	0. 25
	HEMBA1005497	F-HEMBA1005497	0.09
35	HEMBA1005500	F-HEMBA1005500	0.08
	HEMBA1005506	F-HEMBA1005506	0. 17
	HEMBA1005508	F-HEMBA1005508	0.12
•	HEMBA1005511	F-HEMBA1005511	0. 1
40	HEMBA1005520	F-HEMBA1005520	0. 07
	HEMBA1005552	F-HEMBA1005552	0. 05
	HEMBA1005568	F-HEMBA1005568	0.06
	HEMBA1005570	F-HEMBA1005570	0. 1
45	HEMBA1005577	F-HEMBA1005577	0.13
	HEMBA1005588	F-HEMBA1005588	0. 25
•	HEMBA1005593	F-HEMBA1005593	0.13
	HEMBA1005606	F-HEMBA1005606	0.16
50	HEMBA1005616	F-HEMBA1005616	0. 15
	HEMBA1005627	F-HEMBA1005627	0, 11
	HEMBA1005631	F-HEMBA1005631	0.07
	HEMBA1005632	F-HEMBA1005632	0. 07
55	HEMBA1005634	F-HEMBA1005634	0. 07
55	HEMBA1005670	F-HEMBA1005670	0.11

	HEMBA1005679	F-HEMBA1005679	0. 22
	HEMBA1005699	F-HEMBA1005699	0.08
	HEMBA1005705	F-HEMBA1005705	0.08
5	HEMBA1005717	F-HEMBA1005717	
	HEMBA1005765	F-HEMBA1005765	0.08
	HEMBA1005780	F-HEMBA1005780	0.16
	HEMBA1005829	F-HEMBA1005829	0.09
	HEMBA1005853	F-HEMBA1005853	0.06
	HEMBA1005894	F-HEMBA1005894	0.16
	HEMBA1005909	F-HEMBA1005909	0.13
	HEMBA1005911	F-HEMBA1005911	0. 22
15	HEMBA1005921	F-HEMBA1005921	0. 1
	HEMBA1005931	F-HEMBA1005931	0. 16
	HEMBA1005934	F-HEMBA1005934,	
	HEMBA1005962	F-HEMBA1005962	0. 07
	HEMBA1005999	F-HEMBA1005999	0. 14
20	HEMBA1006002	F-HEMBA1006002	0.09
	HEMBA1006005	F-HEMBA1006005	0. 2
	HEMBA1006036	F-HEMBA1006036	0. 24
	HEMBA1006042	F-HEMBA1006042	0.11
25	HEMBA1006067	F-HEMBA1006067	0. 09
	HEMBA1006090	F-HEMBA1006090	0.09
	HEMBA1006124	F-HEMBA1006124	0.05
	HEMBA1006142	F-HEMBA1006142	0. 13
30	HEMBA1006158	F-HEMBA1006158	0. 27
	HEMBA1006253	F-HEMBA1006253	0. 29
	HEMBA1006259	F-HEMBA1006259	0. 11
	HEMBA1006268	F-HEMBA1006268	0.11
35	HEMBA1006328	F-HEMBA1006328	0.06
	HEMBA1006334	F-HEMBA1006334	0.08
	HEMBA1006359	F-HEMBA1006359	0. 22
	HEMBA1006364	F-HEMBA1006364	0. 23
40	HEMBA1006380	F-HEMBA1006380	0.09
	HEMBA1006416	F-HEMBA1006416	0. 25
	HEMBA1006419	F-HEMBA1006419	0.08
	HEMBA1006421	F-HEMBA1006421	0.26
45	HEMBA1006426	F-HEMBA1006426	0.17
45	HEMBA1006438	F-HEMBA1006438	0.08
	HEMBA1006446	F-HEMBA1006446	0.11
	HEMBA1006461	F-HEMBA1006461	0.15
50	HEMBA1006471	F-HEMBA1006471	0.06
50	HEMBA1006486	F-HEMBA1006486	0.11
	HEMBA1006489	F-HEMBA1006489	0.09
	HEMBA1006540	F-HEMBA1006540	0. 23
	HEMBA1006546	F-HEMBA1006546	0.07
55	HEMBA1006562	F-HEMBA1006562	0.12

	HEMBA1006569	F-HEMBA1006569	
	HEMBA1006595	F-HEMBA1006595	0.08
	HEMBA1006597	F-HEMBA1006597	0. 17
5	HEMBA1006617	F-HEMBA1006617	0.07
	HEMBA1006635	F-HEMBA1006635	0.07
	HEMBA1006639	F-HEMBA1006639	0. 25
	HEMBA1006648	F-HEMBA1006648	0. 24
10	HEMBA1006653	F-HEMBA1006653	0.09
	HEMBA1006665	F-HEMBA1006665	0. 17
	HEMBA1006676	F-HEMBA1006676	0.17
	HEMBA1006695	F-HEMBA1006695	0. 1
15	HEMBA1006696	F-HEMBA1006696	0.05
	HEMBA1006744	F-HEMBA1006744	0.12
	HEMBA1006779	F-HEMBA1006779,	
,	HEMBA1006780	F-HEMBA1006780	0. 23
00	HEMBA1006821	F-HEMBA1006821	0. 08
20	HEMBA1006824	F-HEMBA1006824	0. 09
	HEMBA1006849	F-HEMBA1006849	0. 05
	HEMBA1006865	F-HEMBA1006865	0. 1
	HEMBA1006921	F-HEMBA1006921	0, 22
25	HEMBA1006929	F-HEMBA1006929	0. 15
	HEMBA1006938	F-HEMBA1006938	0. 07
	HEMBA1006949	F~HEMBA1006949	0. 27
	HEMBA1007017	F-HEMBA1007017	0. 07
30	HEMBA1007045	F~HEMBA1007045	0.14
	HEMBA1007051	F~HEMBA1007051	0. 19
•	HEMBA1007073	F~HEMBA1007073	0. 22
	HEMBA1007078	F-HEMBA1007078	0.12
35	HEMBA1007113	F-HEMBA1007113	0.12
	HEMBA1007129	F-HEMBA1007129	0. 18
•	HEMBA1007147	F-HEMBA1007147	0.08
	HEMBA1007206	F-HEMBA1007206	0.11
40	HEMBA1007256	F-HEMBA1007256	0.13
	HEMBA1007273	F-HEMBA1007273	0.05
	HEMBA1007279	F-HEMBA1007279	0.09
	HEMBA1007288	F-HEMBA1007288	0.06
45	HEMBA1007322	F-HEMBA1007322	0.12
	HEMBA1007327	F-HEMBA1007327	0.12
	HEMBA1007341	F-HEMBA1007341	0.1
	HEMBA1007347	F-HEMBA1007347	0.14
50	HEMBB1000005	F-HEMBB1000005	0.13
50	HEMBB1000039	F-HEMBB1000039	0.07
	HEMBB1000044	F-HEMB81000044	0.08
	HEMBB1000050	F-HEMB81000050	0. 07
	HEMBB1000054	F-HEMBB1000054	0.12
55	HEMBB1000055	F-HEMBB1000055	0. 2

	HEMBB1000059	F-HEMBB1000059	0. 08
	HEMBB1000089	F-HEMB81000089	0.09
	HEMBB1000099	F-HEMBB1000099	0.17
5	HEMBB1000113	F-HEMBB1000113	0.12
	HEMBB1000141	F-HEMBB1000141	0.13
	HEMBB1000144	F-HEMBB1000144	0. 2
	HEMBB1000218	F-HEMBB1000218	0.07
10	HEMBB1000250	F-HEMBB1000250	0.26
	HEMBB1000258	F-HEMBB1000258	0.15
	HEMBB1000264	F-HEMBB1000264	0.08
	HEMBB1000272	F-HEMBB1000272	0.11
15	HEMBB1000274	F-HEMBB1000274	0.12
	HEMBB1000284	F-HEMBB1000284	0.1
	HEMBB1000307	F-HEMBB1000307,	0.11
	HEMBB1000312	F-HEMBB1000312	0.07
20	HEMBB1000317	F-HEMBB1000317	0.27
20	HEMBB1000318	F-HEMBB1000318	0.11
	HEMBB1000335	F-HEMBB1000335	0.15
	HEMBB1000336	F-HEMBB1000336	0. 28
	HEMBB1000337	F-HEMBB1000337	0. 25
25	HEMBB1000341	F-HEMBB1000341	0. 24
	HEMBB1000343	F-HEMBB1000343	0. 07
	HEMBB1000354	F-HEMBB1000354	0. 15
	HEMBB1000369	F-HEMBB1000369	0.08
30	HEMB81000374	F-HEMBB1000374	0. 3
	HEMB81000376	F-HEMBB1000376	0. 12
	HEMB81000399	F-HEMBB1000399	0.08
	HEMBB1000402	F-HEMBB1000402	0. 29
35	HEMBB1000404	F-HEMBB1000404	0. 29
	HEMBB1000420	F-HEMBB1000420	0.11
	HEMBB1000434	F-HEMB81000434	0.05
	HEMBB1000441	F-HEMBB1000441	0. 12
40	HEMBB1000455	F-HEMBB1000455	0.05
	HEMBB1000472		0. 05 0. 13
	HEMBB1000480	F-HEMBB1000480	
	HEMBB1000487	F-HEMBB1000487	0.09
45	HEMBB1000490	F-HEMBB1000490	
	HEMBB1000491	F-HEMBB1000491 F-HEMBB1000493	0. 16 0. 15
	HEMBB1000493		0. 15
	HEMBB1000518 HEMBB1000523	F-HEMBB1000518 F-HEMBB1000523	0.03
50	HEMBB1000530		0. 07
	HEMBB1000554		0.14
·	HEMBB1000564		0. 09
	HEMBB1000573		0. 03
55	HEMBB1000575	F-HEMBB1000575	0.08
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	HEMBB1000586	F-HEMBB1000586	0.12
	HEMBB1000598	F-HEMBB1000598	0.12
	HEMBB1000637	F-HEMBB1000637	0.19
5	HEMBB1000638	F-HEMBB1000638	0.07
	HEMBB1000643	F-HEMBB1000643	0.07
	HEMBB1000649	F-HEMBB1000649	0.13
	HEMBB1000652	F-HEMBB1000652	0. 2
10	HEMBB1000665	F-HEMBB1000665	0.07
	HEMBB1000684	F-HEMBB1000684	0.06
	HEMBB1000709	F-HEMBB1000709	0.09
	HEMBB1000726	F-HEMBB1000726	0.07
15	HEMBB1000738	F-HEMBB1000738	0.19
	HEMBB1000749	F-HEMBB1000749	0.14
	HEMBB1000770	F-HEMBB1000770,	0.09
	HEMBB1000790	F-HEMBB1000790	0. 23
	HEMBB1000794	F-HEMBB1000794	0.09
20	HEMBB1000821	F-HEMB81000821	0.1
	HEMBB1000822	F-HEMB81000822	0.18
	HEMBB1000827	F-HEMBB1000827	0.1
	HEMBB1000831	F-HEMBB1000831	0. 2
25	HEMBB1000840		0.09
	HEMBB1000876		0.12
	HEMBB1000883		0.13
	HEMBB1000888		0.1
30	HEMBB1000890		0.09
	HEMBB1000893		0.1
	HEMBB1000910		0.06
	HEMBB1000913		0.09
35	HEMBB1000915		0.12
	HEMBB1000917		0. 15
	HEMBB1000959		0.08
	HEMBB1000981		0.09
40	HEMBB1000996		0.1
	HEMBB1001004		0.08
	HEMBB1001008		0. 25
	HEMBB1001037		0.14
45	HEMBB1001047		0.14
	HEMBB1001051		0.13
	HEMBB1001060		0.16
	HEMB81001063		0. 1 0. 19
50	HEMBB1001102		0. 19
	HEMBB1001114 HEMBB1001119		0. 08
			0.13
	HEMBB1001133 HEMBB1001143		0.17
55	HEMBB100117		0. 17
35	TICHIDD FOOTT)	, , ,,_,,_,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	

	HEMBB1001208	F-HEMBB1001208	0. 12
	HEMBB1001209	F-HEMBB1001209	0.11
	HEMBB1001218	F-HEMBB1001218	0.1
5	HEMBB1001221	F-HEMBB1001221	0.1
	HEMBB1001249	F-HEMBB1001249	0.12
	HEMBB1001253	F-HEMB81001253	0.13
	HEMBB1001254	F-HEMBB1001254	0.13
10	HEMBB1001267	F-HEMBB1001267	0.09
	HEMBB1001271	F-HEMB81001271	0.16
	HEMBB1001282	F-HEMBB1001282	0.14
	HEMBB1001302	F-HEMBB1001302	0. 18
15	HEMBB1001304	F-HEMBB1001304	0.17
	HEMBB1001315	F-HEMBB1001315	0.18
	HEMBB1001317	F-HEMBB1001317,	0. 12
	HEMBB1001326	F-HEMBB1001326	0.09
20	HEMBB1001335	F-HEMBB1001335	0.08
	HEMBB1001337	F-HEMBB1001337	0. 12
	HEMBB1001348	F-HEMBB1001348	0. 11
	HEMBB1001356	F-HEMBB1001356	0.1
25	HEMBB1001366	F-HEMBB1001366	0. 16
23	HEMBB1001367	F-HEMB81001367	0. 11
	HEMBB1001380	F-HEMBB1001380	0. 07
	HEMBB1001424	F-HEMBB1001424	0. 05
	HEMBB1001436	F-HEMBB1001436	0. 14
30	HEMBB1001443	F-HEMBB1001443	0. 28
	HEMBB1001454	F-HEMBB1001454	
	HEMBB10C1458	F-HEMBB1001458	0. 11
	HEMBB1001463	F-HEMBB1001463	0. 1
35	HEMBB1001464	F-HEMBB1001464	0.06
	HEMBB1001500	F-HEMBB1001500	0.11
	HEMBB1001521	F-HEMBB1001521	0. 13
	HEMBB1001527	F-HEMBB1001527	0.13
40	HEMBB1001535	F-HEMBB1001535 F-HEMBB1001536	0.11
	HEMBB1001536	F-HEMBB1001537	0.08
	HEMBB1001537	,	
	HEMBB1001555	F-HEMBB1001555	0. 08 0. 1
45	HEMBB1001565	F-HEMBB1001565	
	HEMBB1001586	F-HEMBB1001586 F-HEMBB1001588	0.06 0.06
	HEMBB1001588	F-HEMBB1001618	0.09
	HEMBB1001618 HEMBB1001619	F-HEMBB1001619	0. 09
50	HEMBB1001630	F-HEMBB1001630	0. 12
	HEMBB1001637	F-HEMBB1001637	0. 12
	HEMBB1001641	F-HEMB81001641	0.09
	HEMBB1001704	F-HEMBB1001704	0.09
55	HEMBB1001704	F-HEMBB1001704	0. 11
	11CHDD 1001700	LMDD1001700	V. 11

	HEMBB1001717	F-HEMBB1001717	0.09
	HEMBB1001735	F-HEMBB1001735	0. 21
	HEMBB1001747	F-HEMBB1001747	0.07
5	HEMBB1001749	F-HEMBB1001749	0.13
	HEMBB1001753	F-HEMBB1001753	0.08
	HEMBB1001756	F-HEMBB1001756	0.11
	HEMBB1001762	F-HEMBB1001762	0.11
10	HEMBB1001797	F-HEMBB1001797	0.13
	HEMBB1001802	F-HEMBB1001802	0.06
	HEMBB1001836	F-HEMBB1001836	0.15
	HEMBB1001850	F-HEMBB1001850	0.06
15	HEMBB1001863	F-HEMBB1001863	0.09
	HEMBB1001867	F-HEMBB1001867	0.11
	HEMBB1001868	F-HEMBB1001868,	0.24
	HEMBB1001875	F-HEMBB1001875	0.05
20	HEMBB1001880	F-HEMBB1001880	0.08
	HEMBB1001899	F-HEMBB1001899	0.16
	HEMBB1001911	F-HEMBB1001911	0.22
	HEMBB1001921	F-HEMBB1001921	0.07
25 ·	HEMBB1001922	F-HEMBB1001922	0.11
23	HEMBB1001925	F-HEMBB1001925	0.19
	HEMBB1001930	F-HEMBB1001930	0.1
	HEMBB1001944	F-HEMBB1001944	0.14
20	HEMBB1001952	F-HEMBB1001952	0.08
30	HEMBB1001953	F-HEMBB1001953	0. 15
	HEMBB1001967	F-HEMBB1001967	0.08
	HEMBB1001973	F-HEMBB1001973	0.09
	HEMBB1001983	F-HEMBB1001983	0. 1
35	HEMBB1001988	F-HEMBB1001988	0.15
	HEMBB1001996	F-HEMBB1001996	0.19
	HEMBB1001997	F-HEMBB1001997	0.08
	HEMBB1002002	F-HEMBB1002002	0. 21
40	HEMBB1002005	F-HEMBB1002005	0.18
	HEMB81002009	F-HEMBB1002009	0.09
	HEMBB1002015	F-HEMBB1002015 F-HEMBB1002042	0. 26
	HEMBB1002042 HEMBB1002043	F-HEMBB1002043	0.11
45			0.09
	HEMBB1002045	F-HEMBB1002045	0. 26 0. 12
	HEMBB1002049 HEMBB1002069	F-HEMBB1002049 F-HEMBB1002069	0. 12
	HEMBB1002092	F-HEMBB1002099	0. 12
50	HEMBB1002094	F-HEMBB1002092	0.10
	HEMBB1002094	F-HEMBB1002034	0.09
	HEMBB1002113	F-HEMBB1002139	0. 03
	HEMBB1002189	F-HEMBB1002189	0.06
55	HEMBB1002190	F-HEMBB1002190	0.18
			• •

	HEMBB1002232	F-HEMBB1002232	0. 09
	HEMBB1002247	F-HEMBB1002247	0.17
	HEMBB1002254	F-HEMBB1002254	0.05
5	HEMBB1002255	F-HEMBB1002255	0.11
	HEMBB1002266	F-HEMBB1002266	0. 28
	HEMBB1002280	F-HEMBB1002280	0.07
	HEMBB1002306	F-HEMBB1002306	0.06
10	HEMBB1002364	F-HEMBB1002364	0.05
	HEMBB1002371	F-HEMBB1002371	0.12
	HEMBB1002381	F-HEMBB1002381	0.09
	HEMBB1002387	F-HEMBB1002387	0.17
15	HEMBB1002425	F-HEMBB1002425	0.08
	HEMBB1002453	F-HEMBB1002453	0.12
	HEMB81002458	F-HEMBB1002458,	0.09
	HEMBB1002477	F-HEMBB1002477	0.17
00	HEMBB1002509	F-HEMBB1002509	0.06
20	HEMBB1002520	F-HEMBB1002520	0.08
	HEMBB1002522	F-HEMBB1002522	0.21
	HEMBB1002531	F-HEMBB1002531	0.1
	HEMBB1002534	F-HEMBB1002534	0.09
25	HEMBB1002556	F-HEMBB1002556	0.1
	HEMBB1002579	F-HEMBB1002579	0.13
	HEMBB1002582	F-HEMBB1002582	0. 17
	HEMBB1002590	F-HEMBB1002590	0.08
30	HEMBB1002596	F-HEMBB1002596	0. 15
	HEMBB1002601	F-HEMBB1002601	0. 1
	HEMBB1002603	F-HEMBB1002603	0. 12
	HEMBB1002610	F-HEMBB1002610	0.1
35	HEMBB1002613	F-HEMBB1002613	0.09
	HEMBB1002617	F-HEMBB1002617	0. 12
	HEMBB1002623	F-HEMBB1002623	0. 14
	HEMBB1002635	F-HEMBB1002635	0.08
40	HEMBB1002664	F-HEMB81002664	0.06
	HEMBB1002677	F-HEMBB1002677	0. 07
	HEMBB1002683	F~HEMBB1002683	0.06
	HEMBB1002686	F-HEMBB1002686	0. 12
45	HEMBB1002699	F-HEMBB1002699	0. 15
	HEMBB1002702	F-HEMBB1002702	0.16
	HEMBB1002712	F-HEMBB1002712	0.07
	MAMMA1000009	F-MAMMA1000009	0.11
50	MAMMA1000043	F-MAMMA1000043	0. 19
	MAMMA1000045	F-MAMMA1000045 F-MAMMA1000092	0. 08 0. 11
	MAMMA1000092		0. 11
	MAMMA1000103 MAMMA1000117		0.09
55	MAMMA1000117	F-MAMMA1000117	0. 2
	munimun 1000 129	1 - INCHIBITATION 129	0.17

	MAMMA1000133	F-MAMMA1000133	0. 1
	MAMMA1000134	F-MAMMA1000134	0.05
	MAMMA1000155	F-MAMMA1000155	0. 12
5	MAMMA1000175	F-MAMMA1000175	0. 07
	MAMMA1000198	F-MAMMA1000198	0. 2
	MAMMA1000221	F-MAMMA1000221	0. 1
	MAMMA1000227	F-MAMMA1000227	0.08
10	MAMMA1000241	F-MAMMA1000241	0.16
	MAMMA1000254	F-MAMMA1000254	0.08
	MAMMA1000264	F-MAMMA1000264	0.1
	MAMMA1000266	F-MAMMA1000266	0.18
15	MAMMA1000270	F-MAMMA1000270	0.1
	MAMMA1000287	F-MAMMA1000287	0. 1
	MAMMA1000302	F-MAMMA1000302,	0.07
	MAMMA1000307	F-MAMMA1000307	0.26
20	MAMMA1000331	F-MAMMA1000331	0.18
20	MAMMA1000340	F-MAMMA1000340	0. 2
	MAMMA1000348	F-MAMMA1000348	0.09
	MAMMA1000356	F-MAMMA1000356	0.15
	MAMMA1000360	F-MAMMA1000360	0.09
25	MAMMA1000385	F-MAMMA1000385	0.08
	MAMMA1000402	F-MAMMA1000402	0.11
	MAMMA1000413	F-MAMMA1000413	0. 1
	MAMMA1000414	F-MAMMA1000414	0. 1
30	MAMMA1000423	F-MAMMA1000423	0.06
	MAMMA1000424	F-MAMMA1000424	0. 1
	MAMMA1000431	F-MAMMA1000431	0. 28
	MAMMA1000444	F-MAMMA1000444	0. 07
35	MAMMA1000446	F-MAMMA1000446	0.08
	MAMMA1000478	F-MAMMA1000478	0. 1
	MAMMA1000483	F-MAMMA1000483	0. 07
	MAMMA1000500	F-MAMMA1000500	0. 16
40	MAMMA1000501	F-MAMMA1000501	0.05
	MAMMA1000516	F-MAMMA1000516	0.06
	MAMMA1000522	F-MAMMA1000522	0. 21
	MAMMA1000559	F-MAMMA1000559	0. 12
45	MAMMA1000565	F-MAMMA1000565	0.09
	MAMMA1000576	F-MAMMA1000576	0.14
	MAMMA1000585	F-MAMMA1000585	0.07
	MAMMA1000594	F-MAMMA1000594	0.17
50	MAMMA1000597 MAMMA1000605	F-MAMMA1000597 F-MAMMA1000605	0. 1 0. 09
	MAMMA1000605	F-MAMMA1000616	0.09
	MAMMA1000618	F-MAMMA1000610	0.08
	MAMMA1000621	F-MAMMA1000643	0. 07
55	MAMMA1000649	F-MAMMA1000649	0. 11
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	MAMMA1000696	F-MAMMA1000696	0.08
	MAMMA1000707	F-MAMMA1000707	0.07
	MAMMA1000714	F-MAMMA1000714	0.18
5	MAMMA1000718	F-MAMMA1000718	0.06
	MAMMA1000720	F-MAMMA1000720	0. 15
	MAMMA1000723	F-MAMMA1000723	0.14
	MAMMA1000732	F-MAMMA1000732	0.06
10	MAMMA1000733	F-MAMMA1000733	0.1
	MAMMA1000744	F-MAMMA1000744	0.15
	MAMMA1000752	F-MAMMA1000752	0.12
	MAMMA1000761	F-MAMMA1000761	0.24
15	MAMMA1000775	F-MAMMA1000775	0. 24
	MAMMA1000778	F-MAMMA1000778	0.09
	MAMMA1000782	F-MAMMA1000782,	0.1
	MAMMA1000798	F-MAMMA1000798	0. 16
20	MAMMA1000802	F-MAMMA1000802	0.11
	MAMMA1000839	F-MAMMA1000839	0.09
	MAMMA1000845	F-MAMMA1000845	0.1
	MAMMA1000851	F-MAMMA1000851	0.08
25	MAMMA1000855	F-MAMMA1000855	0. 23
20	MAMMA1000862	F-MAMMA1000862	0.12
	MAMMA1000863	F-MAMMA1000863	0. 16
	MAMMA1000867	F-MAMMA1000867	0. 25
22	MAMMA1000876	F-MAMMA1000876	0.14
30	MAMMA1000877	F-MAMMA1000877	0.07
	MAMMA1000880	F-MAMMA1000880	0. 13
	MAMMA1000883	F-MAMMA1000883	0.3
	MAMMA1000905	F-MAMMA1000905	0. 08 0. 18
35	MAMMA1000931 MAMMA1000940	F-MAMMA1000931 F-MAMMA1000940	0. 18
	MAMMA1000940	F-MAMMA1000940	0. 21
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40	MAMMA1000957	F-MAMMA1000957	0. 13
	MAMMA1000962	F-MAMMA1000962	0.06
	MAMMA1000975	F-MAMMA1000975	0, 26
	MAMMA1000987	F-MAMMA1000987	0.08
45	MAMMA1000998	F-MAMMA1000998	0.08
	MAMMA1001003	F-MAMMA1001003	0. 15
	MAMMA1001024	F-MAMMA1001024	0. 08
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50	MAMMA1001035	F-MAMMA1001035	0.08
	MAMMA1001038	F-MAMMA1001038	0. 15
	MAMMA1001050	F-MAMMA1001050	0.06
	MAMMA1001067	F-MAMMA1001067	0.08
55	MAMMA1001074	F-MAMMA1001074	0.09

	MAMMA1001082	F-MAMMA1001082	0.1
	MAMMA1001092	F-MAMMA1001092	0.05
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5	MAMMA1001145	F-MAMMA1001145	0, 05
	MAMMA1001161	F-MAMMA1001161	0.11
	MAMMA1001162	F-MAMMA1001162	0.11
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10	MAMMA1001191	F-MAMMA1001191	0.08
	MAMMA1001203	F-MAMMA1001203	0.08
	MAMMA1001206	F-MAMMA1001206	0.09
·	MAMMA1001220	F-MAMMA1001220	0.1
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	MAMMA1001249	F-MAMMA1001249	0.13
	MAMMA1001256	F-MAMMA1001256,	0.1
	MAMMA1001268	F-MAMMA1001268	0. 26
	MAMMA1001274	F-MAMMA1001274	0.13
20	MAMMA1001280	F-MAMMA1001280	0.1
	MAMMA1001298	F-MAMMA1001298	0.07
	MAMMA1001322	F-MAMMA1001322	0.18
	MAMMA1001324	F-MAMMA1001324	0. 25
25	MAMMA1001330	F-MAMMA1001330	0. 13
	MAMMA1001341	F-MAMMA1001341	0.16
	MAMMA1001383	F-MAMMA1001383	0.05
	MAMMA1001397	F-MAMMA1001397	0. 22
30	MAMMA1001408	F-MAMMA1001408	0.06
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	MAMMA1001442	F-MAMMA1001442	0.13
35	MAMMA1001446	F-MAMMA1001446	0.08
	MAMMA1001452	F-MAMMA1001452	0.1
	MAMMA1001501	F-MAMMA1001501	0.12
	MAMMA1001502	F-MAMMA1001502 F-MAMMA1001547	0. 09 0. 22
40	MAMMA1001547 MAMMA1001575	F-MAMMA1001547	0. 22
	MAMMA1001590	F-MAMMA1001579	0.14
	MAMMA1001696	F-MAMMA1001606	0.14
	MAMMA1001663		
45	MAMMA1001670	F-MAMMA1001670	0. 2
	MAMMA1001670	F-MAMMA1001671	0.13
	MAMMA1001671	F-MAMMA1001679	0.13
	MAMMA1001683	F-MAMMA1001683	0.16
50	MAMMA1001686	F-MAMMA1001686	0.09
	MAMMA1001711	F-MAMMA1001711	0.11
	MAMMA1001715	F-MAMMA1001715	0.11
	MAMMA1001740		0. 1
55	MAMMA1001745	F-MAMMA1001745	0.09
* *			

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	MAMMA1001769	F-MAMMA1001769	0.09
	MAMMA1001788	F-MAMMA1001788	0.1
5	MAMMA1001806	F-MAMMA1001806	0.06
·	MAMMA1001815	F-MAMMA1001815	0.1
	MAMMA1001818	F-MAMMA1001818	0. 1
	MAMMA1001820	F-MAMMA1001820	0. 27
10	MAMMA1001836	F-MAMMA1001836	0.05
	MAMMA1001854	F-MAMMA1001854	0.17
	MAMMA1001878	F-MAMMA1001878	0. 13
	MAMMA1001880	F-MAMMA1001880	0.15
15	MAMMA1001890	F-MAMMA1001890	0.05
	MAMMA1001907	F-MAMMA1001907	0. 1
	MAMMA1001908	F-MAMMA1001908,	0.3
	MAMMA1001931	F-MAMMA1001931	0.14
20	MAMMA1001963	F-MAMMA1001963	0.08
	MAMMA1001992	F-MAMMA1001992	0.12
	MAMMA1002032	F-MAMMA1002032	0.08
	MAMMA1002056	F-MAMMA1002056	0.09
25	MAMMA1002058	F-MAMMA1002058	0.16
	MAMMA1002078	F-MAMMA1002078	0. 28
	MAMMA1002082	F-MAMMA1002082	0. 08
	MAMMA1002084	F-MAMMA1002084	0. 24
30	MAMMA1002093	F-MAMMA1002093	0. 12
	MAMMA1002108	F-MAMMA1002108	0.05
	MAMMA1002118	F-MAMMA1002118	0. 1
	MAMMA1002125	F-MAMMA1002125 F-MAMMA1002132	0.09
35	MAMMA1002132 MAMMA1002140	F-MAMMA1002140	0. 11
33	MAMMA1002145	F-MAMMA1002145	0. 11
	MAMMA1002145	F-MAMMA1002155	0. 1
	MAMMA1002158	F-MAMMA1002158	0. 11
40	MAMMA1002215	F-MAMMA1002215	0. 05
40	MAMMA1002230	F-MAMMA1002230	0. 07
	MAMMA1002250	F-MAMMA1002250	0. 2
	MAMMA1002267	F-MAMMA1002267	0.08
-	MAMMA1002282	F-MAMMA1002282	0. 05
45	MAMMA1002293	F-MAMMA1002293	0.09
	MAMMA1002298	F-MAMMA1002298	0.09
	MAMMA1002299	F-MAMMA1002299	0.16
50	MAMMA1002310	F-MAMMA1002310	0. 16
50	MAMMA1002311	F-MAMMA1002311	0. 13
	MAMMA1002322	F-MAMMA1002322	0.07
	MAMMA1002332	F-MAMMA1002332	0.08
	MAMMA1002339	F-MAMMA1002339	0. 09
55	MAMMA1002347	F-MAMMA1002347	0.06

	MAMMA1002352	F-MAMMA1002352	0.08
	MAMMA1002359	F-MAMMA1002359	0.13
	MAMMA1002360	F-MAMMA1002360	0.17
5	MAMMA1002361	F-MAMMA1002361	0.13
	MAMMA1002392	F-MAMMA1002392	0.16
	MAMMA1002411	F-MAMMA1002411	0.28
	MAMMA1002417	F-MAMMA1002417	0.19
10	MAMMA1002428	F-MAMMA1002428	0.07
	MAMMA1002446	F-MAMMA1002446	0.17
	MAMMA1002475	F-MAMMA1002475	0.17
	MAMMA1002480	F-MAMMA1002480	0.07
15	MAMMA1002494	F-MAMMA1002494	0.17
	MAMMA1002498	F-MAMMA1002498	0. 28
	MAMMA1002545	F-MAMMA1002545,	0.12
	MAMMA1002556	F-MAMMA1002556	0.06
20	MAMMA1002566	F-MAMMA1002566	0.13
	MAMMA1002571	F-MAMMA1002571	0.29
	MAMMA1002573	F-MAMMA1002573	0. 29
	MAMMA1002597	F-MAMMA1002597	0.1
25	MAMMA1002603	F-MAMMA1002603	0. 1
25	MAMMA1002612	F-MAMMA1002612	0.14
	MAMMA1002618	F-MAMMA1002618	0.11
	MAMMA1002622	F-MAMMA1002622	0. 24
	MAMMA1002623	F-MAMMA1002623	0. 1
30	MAMMA1002625	F-MAMMA1002625	0. 11
	MAMMA1002629	F-MAMMA1002629	0.05
	MAMMA1002636	F-MAMMA1002636	0. 12
	MAMMA1002646	F-MAMMA1002646	0.05
35	MAMMA1002662	F-MAMMA1002662	0.09
	MAMMA1002698	F-MAMMA1002698	0.08
	MAMMA1002701 MAMMA1002708	F-MAMMA1002701 F-MAMMA1002708	0. 11
	MAMMA1002708	F-MAMMA1002708	0.09
40	MAMMA1002727	F-MAMMA1002727	0. 15
	MAMMA1002727	F-MAMMA1002728	0. 13
	MAMMA1002744	F-MAMMA1002744	0. 12
	MAMMA1002746	F-MAMMA1002746	0.09
45	MAMMA1002748	F-MAMMA1002748	0.1
	MAMMA1002754	F-MAMMA1002754	0.07
	MAMMA1002758	F-MAMMA1002758	0. 29
	MAMMA1002764	F-MAMMA1002764	0. 2
50	MAMMA1002765	F-MAMMA1002765	0. 29
	MAMMA1002780	F-MAMMA1002780	0.12
	MAMMA1002820	F-MAMMA1002820	0.3
	MAMMA1002830	F-MAMMA1002830	0.16
55	MAMMA1002833	F-MAMMA1002833	0.14

	MAMMA1002835	F-MAMMA1002835	0. 13
	MAMMA1002838	F-MAMMA1002838	0.09
	MAMMA1002844	F-MAMMA1002844	0. 27
5	MAMMA1002858	F-MAMMA1002858	0. 29
	MAMMA1002871	F-MAMMA1002871	0.11
	MAMMA1002880	F-MAMMA1002880	0.11
	MAMMA1002887	F-MAMMA1002887	0.09
10	MAMMA1002892	F-MAMMA1002892	0. 17
•	MAMMA1002908	F-MAMMA1002908	0.11
	MAMMA1002909	F-MAMMA1002909	0. 22
	MAMMA1002930	F-MAMMA1002930	0. 2
15	MAMMA1002941	F-MAMMA1002941	0. 23
	MAMMA1002970	F-MAMMA1002970	0.06
	MAMMA1002972	F-MAMMA1002972,	0.08
	MAMMA1002973	F-MAMMA1002973	0.1
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	MAMMA1003003	F-MAMMA1003003	0. 2
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05	MAMMA1003019	F-MAMMA1003019	0.12
25	MAMMA1003026	F-MAMMA1003026	0.26
	MAMMA1003031	F-MAMMA1003031	0.18
	MAMMA1003039	F-MAMMA1003039	0.09
	MAMMA1003040	F-MAMMA1003040	0.07
30	MAMMA1003055	F-MAMMA1003055	0.08
	MAMMA1003089	F-MAMMA1003089	0.14
	MAMMA1003140	F-MAMMA1003140	0.19
	NT2RM2000609	F-NT2RM2000609	0. 26
35	NT2RM4000046	F-NT2RM4000046	0. 1
	NT2RM4000085	F-NT2RM4000085	0. 23
	NT2RM4000086	F-NT2RM4000086	0. 28
	NT2RM4000200	F-NT2RM4000200	0.16
40	NT2RM4000244	F-NT2RM4000244	0.07
	NT2RM4000265	F-NT2RM4000265	0.18
	NT2RM4000327	F-NT2RM4000327	0.06
	NT2RM4000366	F-NT2RM4000366	0. 28
45	NT2RM4000368	F-NT2RM4000368	0.11
	NT2RM4000414	F-NT2RM4000414	0.19
	NT2RM4000425	F-NT2RM4000425	0. 22
	NT2RM4000532	F-NT2RM4000532	0.14
50	NT2RM4000779	F-NT2RM4000779	0.13
50	NT2RM4000855	F-NT2RM4000855	0.13
	NT2RM4000979	F-NT2RM4000979	0. 22
	NT2RM4001016	F-NT2RM4001016	0. 2
	NT2RM4001160	F-NT2RM4001160	0.1
55	NT2RM4001313	F-NT2RM4001313	0. 26

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	NT2RM4001414	F-NT2RM4001414	0. 23
	NT2RM4001437	F-NT2RM4001437	0.12
_	NT2RM4001519	F-NT2RM4001519	0.07
5	NT2RM4001557	F-NT2RM4001557	0.3
	NT2RM4001605	F-NT2RM4001605	0. 22
	NT2RM4001754	F-NT2RM4001754	0. 26
	NT2RM4001776	F-NT2RM4001776	0.18
10	NT2RM4001810	F-NT2RM4001810	0. 22
	NT2RM4001856	F-NT2RM4001856	0.11
	NT2RM4001858	F-NT2RM4001858	0.14
	NT2RM4001930	F-NT2RM4001930	0.3
15	NT2RM4001953	F-NT2RM4001953	0.09
	NT2RM4001984	F-NT2RM4001984	0.09
	NT2RM4002067	F-NT2RM4002067,	0.08
	NT2RM4002278	F-NT2RM4002278	0. 07
20	NT2RM4002281	F-NT2RM4002281	0.07
	NT2RM4002287	F-NT2RM4002287	0. 27
	NT2RM4002383	F-NT2RM4002383	0.08
	NT2RM4002390	F-NT2RM4002390	0.18
25	NT2RM4002438	F-NT2RM4002438	0. 23
	NT2RM4002479	F-NT2RM4002479	0.27
	NT2RM4002499	F-NT2RM4002499	0.08
	NT2RM4002504	F-NT2RM4002504	0.08
••	NT2RM4002532	F-NT2RM4002532	0. 15
30	NT2RM4002567	F-NT2RM4002567	0. 28
	NT2RP2000027	F-NT2RP2000027	0.07
	NT2RP2000076	F-NT2RP2000076	0.16
	NT2RP2000077	F-NT2RP2000077	0. 2
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	NT2RP2000108	F-NT2RP2000108	0.07
	NT2RP2000183	F-NT2RP2000183	0.12
	NT2RP2000257	F-NT2RP2000257	0. 27
40	NT2RP2000258	F-NT2RP2000258	0.19
	NT2RP2000270	F-NT2RP2000270	0.06
	NT2RP2000289	F-NT2RP2000289	0.06
	NT2RP2000327	F-NT2RP2000327	0.08
45	NT2RP2000337	F-NT2RP2000337	0.14
	NT2RP2000420	F-NT2RP2000420	0.2
	NT2RP2000459	F-NT2RP2000459	0. 12
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	NT2RP2000603 NT2RP2000644	F-NT2RP2000603 F-NT2RP2000644	0. 22
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	NT2RP2000678	F-NT2RP2000678	0. 11
55	NT2RP2000715	F-NT2RP2000713	0. 21
· ·	41714 TACA TACA	1 MIENIZUUU/31	0.09

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	NT2RP2000842	F-NT2RP2000842	0.14
	NT2RP2000970	F-NT2RP2000970	0. 12
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	NT2RP2001149	F-NT2RP2001149	0.08
	NT2RP2001196	F-NT2RP2001196	0. 27
	NT2RP2001226	F-NT2RP2001226	0.18
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	NT2RP2001295	F-NT2RP2001295	0.13
	NT2RP2001312	F-NT2RP2001312	0. 15
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	NT2RP2001423	F-NT2RP2001423	0.06
	NT2RP2001445	F-NT2RP2001445,	
	NT2RP2001449	F-NT2RP2001449	0.15
20	NT2RP2001467	F-NT2RP2001467	0. 21
20	NT2RP2001506	F-NT2RP2001506	0.06
	NT2RP2001526	F-NT2RP2001526	0. 1
	NT2RP2001569	F-NT2RP2001569	0.09
	NT2RP2001663	F-NT2RP2001663	0.08
25	NT2RP2001677	F-NT2RP2001677	0.14
	NT2RP2001678	F-NT2RP2001678	0.17
	NT2RP2001699	F-NT2RP2001699	0.05
	NT2RP2001720	F-NT2RP2001720	0.08
30	NT2RP2001740	F-NT2RP2001740	0. 23
	NT2RP2001762	F-NT2RP2001762	0.11
	NT2RP2001813	F-NT2RP2001813	0.19
	NT2RP2001869	F-NT2RP2001869	0. 22
35	NT2RP2001926	F-NT2RP2001926	0.07
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	NT2RP2002172	F-NT2RP2002172	0. 21
50	NT2RP2002192	F-NT2RP2002192	0. 05
50	NT2RP2002219	F-NT2RP2002219	0. 13
	NT2RP2002256	F-NT2RP2002256	0.17
	NT2RP2002259	F-NT2RP2002259	0. 28
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55	NT2RP2002394	F-NT2RP2002394	0.16

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	NT2RP2002457	F-NT2RP2002457	0.11
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5	NT2RP2002504	F-NT2RP2002504	0. 21
	NT2RP2002546	F-NT2RP2002546	0.17
	NT2RP2002591	F-NT2RP2002591	0.15
	NT2RP2002606	F-NT2RP2002606	0.11
10	NT2RP2002643	F-NT2RP2002643	0.26
	NT2RP2002727	F-NT2RP2002727	0.26
	NT2RP2002736	F-NT2RP2002736	0.12
	NT2RP2002740	F-NT2RP2002740	0.16
16	NT2RP2002741	F-NT2RP2002741	0.11
	NT2RP2002750	F-NT2RP2002750	0.14
	NT2RP2002752	F-NT2RP2002752	0.1
	NT2RP2002753	F-NT2RP2002753	0.09
	NT2RP2002778	F-NT2RP2002778	0.13
20	NT2RP2002839	F-NT2RP2002839	
	NT2RP2002857	F-NT2RP2002857	0. 21
	NT2RP2002987	F-NT2RP2002987	0.08
	NT2RP2003000	F-NT2RP2003000	0. 24
25	NT2RP2003073	F-NT2RP2003073	0.08
	NT2RP2003129	F-NT2RP2003129	0.07
	NT2RP2003161	F-NT2RP2003161	0.05
	NT2RP2003164	F-NT2RP2003164	0.09
30	NT2RP2003206	F-NT2RP2003206	0. 22
	NT2RP2003230	F-NT2RP2003230	0.08
	NT2RP2003237	F-NT2RP2003237	0. 24
	NT2RP2003339	F-NT2RP2003339	0. 1
35	NT2RP2003394	F-NT2RP2003394	0. 07
	NT2RP2003456	F-NT2RP2003456	0.08
	NT2RP2003499	F-NT2RP2003499	0.08
	NT2RP2003517	F-NT2RP2003517	0. 29
40	NT2RP2003522	F-NT2RP2003522	0. 22
40	NT2RP2003559	F-NT2RP2003559	0.12
	NT2RP2003668	F-NT2RP2003668	0.08
	NT2RP2003704	F-NT2RP2003704	0. 09
	NT2RP2003706		0. 28
45	NT2RP2003727	F-NT2RP2003727	0. 17
	NT2RP2003770	F-NT2RP2003770	0.09
	NT2RP2003859	F-NT2RP2003859	0. 18
	NT2RP2003871		0. 23
50	NT2RP2003885	F-NT2RP2003885	0. 24
	NT2RP2003912		0. 20
	NT2RP2003968	F-NT2RP2003968	0. 2
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55	NT2RP2004014	F-NT2RP2004014	0.0

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	NT2RP2004170	F-NT2RP2004170	0.09
	NT2RP2004172	F-NT2RP2004172	0.15
5	NT2RP2004207	F-NT2RP2004207	0.11
	NT2RP2004232	F-NT2RP2004232	0.17
	NT2RP2004270	F-NT2RP2004270	0. 29
	NT2RP2004300	F-NT2RP2004300	0.11
10	NT2RP2004321	F-NT2RP2004321	0.13
	NT2RP2004339	F-NT2RP2004339	0.14
	NT2RP2004347	F-NT2RP2004347	0.14
	NT2RP2004396	F-NT2RP2004396	0. 29
15	NT2RP2004400	F-NT2RP2004400	0.06
	NT2RP2004412	F-NT2RP2004412	0. 24
	NT2RP2004425	F-NT2RP2004425	0.13
	NT2RP2004490	F-NT2RP2004490	0.13
20	NT2RP2004512	F-NT2RP2004512	0.1
•	NT2RP2004523	F-NT2RP2004523	0.15
	NT2RP2004580	F-NT2RP2004580	0.12
	NT2RP2004587	F-NT2RP2004587	0.08
25	NT2RP2004675	F-NT2RP2004675	0.14
	NT2RP2004681	F-NT2RP2004681	0.17
	NT2RP2004709	F-NT2RP2004709	0.08
	NT2RP2004736	F-NT2RP2004736	0.13
20	NT2RP2004767	F-NT2RP2004767	0.21
30	NT2RP2004775	F-NT2RP2004775	0.17
	NT2RP2004961	F-NT2RP2004961	0.21
	NT2RP2004962	F-NT2RP2004962	0.24
	NT2RP2004967	F-NT2RP2004967	0.05
35	NT2RP2004982	F-NT2RP2004982	0.11
	NT2RP2005020	F-NT2RP2005020	0.16
	NT2RP2005031	F-NT2RP2005031	0.07
	NT2RP2005108	F-NT2RP2005108	0.11
40	NT2RP2005254	F-NT2RP2005254	0.16
	NT2RP2005289	F-NT2RP2005289	0. 22
	NT2RP2005293	F-NT2RP2005293	0.08
	NT2RP2005325	F-NT2RP2005325	
45	NT2RP2005336	F-NT2RP2005336	0. 27
	NT2RP2005344	F-NT2RP2005344	0. 24
	NT2RP2005354	F-NT2RP2005354	0.08
	NT2RP2005360	F-NT2RP2005360	0.12
50	NT2RP2005407 NT2RP2005476	F-NT2RP2005407 F-NT2RP2005476	0. 28 0. 14
	NT2RP2005476	F-NT2RP2005476	0. 14
	NT2RP2005491	F-NT2RP2005501	0. 18
	NT2RP2005531	F-NT2RP2005531	0. 1
55	NT2RP2005581	F-NT2RP2005581	0. 23
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	NT2RP2005645	F-NT2RP2005645	0. 13
	NT2RP2005651	F-NT2RP2005651	0.06
	NT2RP2005694	F-NT2RP2005694	0. 3
5	NT2RP2005701	F-NT2RP2005701	0, 11
	NT2RP2005719	F-NT2RP2005719	0.06
	NT2RP2005726	F-NT2RP2005726	0. 29
	NT2RP2005741	F-NT2RP2005741	0.22
10	NT2RP2005753	F-NT2RP2005753	0.16
	NT2RP2005815	F-NT2RP2005815	0.18
	NT2RP2005841	F-NT2RP2005841	0.05
	NT2RP2005857	F-NT2RP2005857	0. 28
15	NT2RP2005908	F-NT2RP2005908	0.05
15	NT2RP2005942	F-NT2RP2005942	0. 26
•	NT2RP2005980	F-NT2RP2005980,	0.1
	NT2RP2006098	F-NT2RP2006098	0.06
	NT2RP2006103	F-NT2RP2006103	0.07
20	NT2RP2006166	F-NT2RP2006166	0. 27
	NT2RP2006184	F-NT2RP2006184	0. 23
	NT2RP2006258	F-NT2RP2006258	0.12
	NT2RP2006261	F-NT2RP2006261	0.16
25	NT2RP2006320	F-NT2RP2006320	0.16
	NT2RP2006321	F-NT2RP2006321	0. 15
·	NT2RP2006323	F-NT2RP2006323	
	NT2RP2006393	F-NT2RP2006393	0. 07
30	NT2RP2006454	F-NT2RP2006454	0.11
	NT2RP2006467	F-NT2RP2006467	0. 24
	NT2RP2006534	F-NT2RP2006534	0.14
	NT2RP2006554	F-NT2RP2006554	0. 16
35	NT2RP2006598	F-NT2RP2006598	0.08
	NT2RP3000055	F-NT2RP3000055	0.14
	NT2RP3000109	F-NT2RP3000109	0.18
	NT2RP3000186	F-NT2RP3000186 F-NT2RP3000233	0. 06 0. 29
40	NT2RP3000233	F-NT2RP3000235	0. 29
	NT2RP3000235	F-NT2RP3000233	0. 19
	NT2RP3000247	F-NT2RP3000247	0. 15
	NT2RP3000341 NT2RP3000418	F-NT2RP3000341	0. 16
45		F-NT2RP3000418	0. 13
	NT2RP3000433 NT2RP3000441	F-NT2RP3000441	0. 11
	NT2RP3000441	F-NT2RP3000449	0.16
	NT2RP3000451	F-NT2RP3000451	0.14
50	NT2RP3000512	F-NT2RP3000512	0. 08
50	NT2RP3000542	F-NT2RP3000542	0. 1
	NT2RP3000561	F-NT2RP3000561	0. 1
	NT2RP3000578		0. 3
	NT2RP3000582		0. 08
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	NT2RP3000584	F-NT2RP3000584	0. 07
	NT2RP3000622	F-NT2RP3000622	0.17
	NT2RP3000628	F-NT2RP3000628	0. 08
5	NT2RP3000685	F-NT2RP3000685	0.17
	NT2RP3000736	F-NT2RP3000736	0. 26
	NT2RP3000742	F-NT2RP3000742	0. 22
	NT2RP3000815	F-NT2RP3000815	0.06
10	NT2RP3000865	F-NT2RP3000865	0.13
	NT2RP3000875	F-NT2RP3000875	0.16
	NT2RP3000904	F-NT2RP3000904	0.13
	NT2RP3001007	F-NT2RP3001007	0.1
15	NT2RP3001115	F-NT2RP3001115	0.1
	NT2RP3001232	F-NT2RP3001232	0.09
	NT2RP3001245	F-NT2RP3001245	0.09
	NT2RP3001281	F-NT2RP3001281	0. 29
20	NT2RP3001318	F-NT2RP3001318	0. 12
	NT2RP3001339	F-NT2RP3001339	0. 27
	NT2RP3001340	F-NT2RP3001340	0.11
	NT2RP3001374	F-NT2RP3001374	0. 21
05	NT2RP3001383	F-NT2RP3001383	0.09
25	NT2RP3001432	F-NT2RP3001432	0, 29
	NT2RP3001459	F-NT2RP3001459	0.16
	NT2RP3001527	F-NT2RP3001527	0.3
	NT2RP3001580	F-NT2RP3001580	0. 23
30	NT2RP3001589	F-NT2RP3001589	0. 12
	NT2RP3001607	F-NT2RP3001607	0.12
	NT2RP3001634	F-NT2RP3001634	0. 15
	NT2RP3001752	F-NT2RP3001752	0.09
35	NT2RP3001782	F-NT2RP3001782	0. 21
	NT2RP3001898	F-NT2RP3001898	0.07
	NT2RP3001926	F-NT2RP3001926	0.05
	NT2RP3001989	F-NT2RP3001989	0.14
40	NT2RP3002002	F-NT2RP3002002	0.18
	NT2RP3002004	F-NT2RP3002004	0. 11 0. 16
	NT2RP3002056	F-NT2RP3002056	0. 16
	NT2RP3002057	F-NT2RP3002057 F-NT2RP3002166	0. 08
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	NT2RP3002007	F-NT2RP3002713	0. 12
50	NT2RP3002713	F-NT2RP3002770	0. 1
	NT2RP3002770	F-NT2RP3002779	0. 23
	NT2RP3002799	F-NT2RP3002810	0. 17
	NT2RP3002910	F-NT2RP3002955	0. 17
55	NT2RP3002938	F-NT2RP3002978	0.06
55	H12M 00028/8	1 H1 ZIII 3002070	0.00

•	NT2RP3003032	F-NT2RP3003032	0. 1
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	NT2RP3003121	F-NT2RP3003121	0. 23
5	NT2RP3003133	F-NT2RP3003133	0. 21
	NT2RP3003138	F-NT2RP3003138	0. 29
	NT2RP3003264	F-NT2RP3003264	0.15
	NT2RP3003346	F-NT2RP3003346	0.16
10	NT2RP3003384	F-NT2RP3003384	0.12
	NT2RP3003403	F-NT2RP3003403	0. 26
	NT2RP3003433	F-NT2RP3003433	0. 22
	NT2RP3003464	F-NT2RP3003464	0. 28
15	NT2RP3003500	F-NT2RP3003500	0. 2
.~	NT2RP3003572	F-NT2RP3003572	0. 29
	NT2RP3003576	F-NT2RP3003576,	
	NT2RP3003625	F-NT2RP3003625	0. 29
	NT2RP3003665	F-NT2RP3003665	0. 15
20	NT2RP3003746	F-NT2RP3003746	0. 07
	NT2RP3003799	F-NT2RP3003799	0.13
	NT2RP3003800	F-NT2RP3003800	0. 28
	NT2RP3003819	F-NT2RP3003819	0. 19
25	NT2RP3003828	F-NT2RP3003828	0.12
•	NT2RP3003842	F-NT2RP3003842	0.11
	NT2RP3003989	F-NT2RP3003989	0.1
	NT2RP3004016	F-NT2RP3004016	0.09
30	NT2RP3004070	F-NT2RP3004070	0.11
	NT2RP3004095	F-NT2RP3004095	0.1
	NT2RP3004110	F-NT2RP3004110	0.07
	NT2RP3004145	F-NT2RP3004145	0.15
35	NT2RP3004148	F-NT2RP3004148	0.14
	NT2RP3004215	F-NT2RP3004215	0.16
	NT2RP3004334	F-NT2RP3004334	0.1
	NT2RP3004349	F-NT2RP3004349	0.09
	NT2RP3004399	F-NT2RP3004399	0.16
40	NT2RP3004470	F-NT2RP3004470	0. 23
	NT2RP3004475	F-NT2RP3004475	0. 26
	NT2RP3004503	F-NT2RP3004503	0.12
	NT2RP3004527	F-NT2RP3004527	0.06
45	NT2RP4000023	F-NT2RP4000023	0.13
	NT2RP4000035	F-NT2RP4000035	0.17
	NT2RP4000049	F-NT2RP4000049	0. 1
	NT2RP4000078	F-NT2RP4000078	0. 28
50	NT2RP4000102	F-NT2RP4000102	0. 15
	NT2RP4000167	F-NT2RP4000167	0. 19
	NT2RP4000214	F-NT2RP4000214	0.12
	NT2RP4000218	F-NT2RP4000218	0.17
55	NT2RP4000321	F-NT2RP4000321	0. 18

	NT2RP4000424	F-NT2RP4000424	0. 22
	NT2RP4000515	F-NT2RP4000515	0.13
5	NT2RP4000517	F-NT2RP4000517	0. 1
5	NT2RP4000519	F-NT2RP4000519	0. 26
	NT2RP4000915	F-NT2RP4000915	0.19
	NT2RP4000996	F-NT2RP4000996	0.05
	NT2RP4001407	F-NT2RP4001407	0. 2
10	NT2RP4001889	F-NT2RP4001889	0. 09
	NT2RP4002075	F-NT2RP4002075	0.19
	NT2RP4002905	F-NT2RP4002905	0.15
	0VARC1000017	F-0VARC1000017	0.12
15	0VARC1000058	F-0VARC1000058	0. 2
	0VARC1000068	F-0VARC1000068	0.14
	0VARC1000071	F-0VARC1000071,	0. 15
	0VARC1000085	F-0VARC1000085	0.16
20	0VARC1000092	F-0VARC1000092	0. 26
	OVARC1000133	F-0VARC1000133	0.14
	0VARC1000145	F-0VARC1000145	0.15
	OVARC1000191	F-0VARC1000191	0.07
25	OVARC1000198	F-0VARC1000198	0. 23
	0VARC1000240	F-0VARC1000240	0.11
	OVARC1000302	F-0VARC1000302	0.08
	0VARC1000414	F-0VARC1000414	0. 12
20	0VARC1000427	F-0VARC1000427	0. 1
30	0VARC1000431	F-0VARC1000431	0.06
	OVARC1000486	F-0VARC1000486	0.08
	0VARC1000496	F-0VARC1000496	0. 13
	0VARC1000526	F-0VARC1000526	0.09
35	0VARC1000533	F-0VARC1000533	0.11
	OVARC1000543	F-0VARC1000543	0. 2
	0VARC1000573	F-0VARC1000573	0.09
	0VARC1000578	F-0VARC1000578	0.13
40	0VARC1000678	F-0VARC1000678	0.17
	0VARC1000679	F-0VARC1000679	0. 23
	0VARC1000700	F-0VARC1000700	0. 28
	0VARC1000769	F-0VARC1000769	0.05
45	0VARC1000802	F-0VARC1000802	0.06
	0VARC1000891		0.08
	0VARC1000897	F-0VARC1000897	0.11
	0VARC1000936	F-0VARC1000936	0.14
50	0VARC1000937	F-0VARC1000937	0. 29
•	0VARC1000948	F-0VARC1000948	0.14
	OVARC1000960	F-0VARC1000960	0.07
	0VARC1000971	F-0VARC1000971	0.11
	0VARC1001000	F-0VARC1001000	0.06
55	OVARC1001011	F-0VARC1001011	0.15

	OVARC1001051	F-0VARC1001051	0. 16
	OVARC1001062	F-0VARC1001062	0.09
	OVARC1001072	F-0VARC1001072	0.11
5	OVARC1001085	F-0VARC1001085	0.12
	OVARC1001117	F-0VARC1001117	0. 21
	OVARC1001118	F-0VARC1001118	0.08
	OVARC1001129	F-0VARC1001129	0. 28
10	OVARC1001169	F-0VARC1001169	0.12
	OVARC1001240	F-0VARC1001240	0.09
	OVARC1001261	F-0VARC1001261	0. 1
	OVARC1001268	F-0VARC1001268	0. 05
15	OVARC1001282	F-0VARC1001282	0.09
	OVARC1001330	F-0VARC1001330	0.08
	OVARC1001339	F-0VARC1001339,	0.15
	OVARC1001342	F-0VARC1001342	0.16
20	OVARC1001357	F-0VARC1001357	0.06
20	OVARC1001442	F-0VARC1001442	0. 15
	OVARC1001480	F-0VARC1001480	0. 23
	OVARC1001542	F-0VARC1001542	0.13
	OVARC1001547	F-0VARC1001547	0.12
25	OVARC1001600	F-0VARC1001600	0.13
	OVARC1001615	F-0VARC1001615	0.11
	OVARC1001668	F-0VARC1001668	0.05
	OVARC1001726	F-0VARC1001726	0.18
30	OVARC1001745	F-0VARC1001745	0. 22
	<ul> <li>OVARC1001795</li> </ul>	F-0VARC1001795	0. 28
	OVARC1001805	F-0VARC1001805	0.16
•	OVARC1001812	F-0VARC1001812	0. 05
35	OVARC1001813	F-0VARC1001813	0. 11
	OVARC1001846	F-0VARC1001846	0. 13
	0VARC1001883	F-0VARC1001883	0. 05
	0VARC1001900	F-0VARC1001900	0. 28
40	0VARC1001901	F-0VARC1001901	0. 09
40	0VARC1001911	F-0VARC1001911	0. 06
	0VARC1001928	F-0VARC1001928	0. 1
	OVARC1001989	F-0VARC1001989	0. 07
	0VARC1002044	F-0VARC1002044	0.06
45	0VARC1002066	F-0VARC1002066	0. 08
	0VARC1002143	F-0VARC1002143	0. 27
	PLACE1000031	F-PLACE1000031	0. 09
	PLACE1000040	F-PLACE1000040	0. 15
50	PLACE1000078	F-PLACE1000078	0. 12
	PLACE1000094	F-PLACE1000094	0.16
	PLACE1000214	F-PLACE1000214	0. 15
	PLACE1000292	F-PLACE1000292	0. 05
	PLACE1000332	F-PLACE1000332	0. 1

	PLACE1000421	F-PLACE1000421	0.14
	PLACE1000424	F-PLACE1000424	0.08
	PLACE1000481	F-PLACE1000481	0.12
5	PLACE1000540	F-PLACE1000540	0.11
	PLACE1000583	F-PLACE1000583	0.1
	PLACE1000599	F-PLACE1000599	0. 22
	PLACE1000749	F-PLACE1000749	0. 2
10	PLACE1000841	F-PLACE1000841	0.09
	PLACE1000931	F-PLACE1000931	0. 22
	PLACE1000979	F-PLACE1000979	0.16
	PLACE1001007	F-PLACE1001007	0.19
15	PLACE1001015	F-PLACE1001015	0. 25
	PLACE1001076	F-PLACE1001076	0. 08
	PLACE1001088	F-PLACE1001088,	0.18
	PLACE1001118	F-PLACE1001118	0. 24
20	PLACE1001136	F-PLACE1001136	0.12
20	PLACE1001272	F-PLACE1001272	0.07
	PLACE1001279	F-PLACE1001279	0.08
	PLACE1001323	F-PLACE1001323	0.18
	PLACE1001377	F-PLACE1001377	0.16
25	PLACE1001384	F-PLACE1001384	0.06
	PLACE1001395	F-PLACE1001395	0.08
	PLACE1001414	F-PLACE1001414	0.11
	PLACE1001440	F-PLACE1001440	0.14
30	PLACE1001468	F-PLACE1001468	0.05
	PLACE1001502	F-PLACE1001502	0.06
	PLACE1001517	F-PLACE1001517	0.09
	PLACE1001534	F-PLACE1001534	0.08
35	PLACE1001603	F-PLACE1001603	0.08
	PLACE1001611	F-PLACE1001611	0. 25
	PLACE1001640	F-PLACE1001640	0.09
	PLACE1001672	F-PLACE1001672	0.11
40	PLACE1001691	F-PLACE1001691	0.11
	PLACE1001705	F-PLACE1001705	0.1
	PLACE1001716	F-PLACE1001716	0.15
	PLACE1001720	F-PLACE1001720	0.18
45	PLACE1001745	F-PLACE1001745	0.09
45	PLACE1001746	F-PLACE1001746	0.12
	PLACE1001748	F-PLACE1001748	0.08
	PLACE1001756	F-PLACE1001756	0.17
	PLACE1001799	F-PLACE1001799	0.14
50	PLACE1001821	F-PLACE1001821	0.09
	PLACE1001845	F-PLACE1001845	0.08
	PLACE1001897	F-PLACE1001897	0.15
	PLACE1002052	F-PLACE1002052	0.13
55	PLACE1002066	F-PLACE1002066	0.07

	PLACE1002150	F-PLACE1002150	0.08
	PLACE1002157	F-PLACE1002157	0.15
	PLACE1002205	F-PLACE1002205	0.13
5	PLACE1002227	F-PLACE1002227	0.12
	PLACE1002256	F-PLACE1002256	0.11
	PLACE1002259	F-PLACE1002259	0.09
•	PLACE1002319	F-PLACE1002319	0.27
10	PLACE1002399	F-PLACE1002399	0.08
	PLACE1002477	F-PLACE1002477	0.17
	PLACE1002493	F-PLACE1002493	0. 09
	PLACE1002500	F-PLACE1002500	0. 25
15	PLACE1002514	F-PLACE1002514	0.06
	PLACE1002537	F-PLACE1002537	0.17
	PLACE1002578	F-PLACE1002578,	
	PLACE1002583	F-PLACE1002583	0. 25
	PLACE1002591	F-PLACE1002591	0. 28
20	PLACE1002598	F-PLACE1002598	0. 29
	PLACE1002604	F-PLACE1002604	0. 15
	PLACE1002768	F-PLACE1002768	0. 11
	PLACE1002772	F-PLACE1002772	0. 15
25	PLACE1002782	F-PLACE1002782	0. 16
	PLACE1002853	F-PLACE1002853	0. 16
	PLACE1002881	F-PLACE1002881	0.09
	PLACE1002962	F-PLACE1002962	0.13
30	PLACE1002968	F-PLACE1002968	0. 23
	PLACE1003153	F-PLACE1003153	0. 07
	PLACE1003205	F-PLACE1003205	0. 2
	PLACE1003238	F-PLACE1003238	0. 22
35	PLACE1003258	F-PLACE1003258	0. 17
	PLACE1003343	F-PLACE1003343	0. 07
	PLACE1003361	F-PLACE1003361	0. 19
	PLACE1003373	F-PLACE1003373	0. 08
40	PLACE1003375	F-PLACE1003375	0. 26
70	PLACE1003401	F-PLACE1003401	0.11
	PLACE1003420	F-PLACE1003420	0. 22
	PLACE1003478	F-PLACE1003478	0. 14
•	PLACE1003516	F-PLACE1003516	0. 17
45	PLACE1003519	F-PLACE1003519	0.14
	PLACE1003521	F-PLACE1003521	0. 17
	PLACE1003528	F-PLACE1003528	0. 16
	PLACE1003566	F-PLACE1003566	0. 16
50	PLACE1003575	F-PLACE1003575	0.06
	PLACE1003583		0.08
	PLACE1003584		0.1
	PLACE1003593	F-PLACE1003593	0.15
55	PLACE1003618	F-PLACE1003618	0. 25

	PLACE1003638	F-PLACE1003638	0.06
	PLACE1003760	F-PLACE1003760	0.12
	PLACE1003768	F-PLACE1003768	0.13
5	PLACE1003833	F-PLACE1003833	0.06
	PLACE1003850	F-PLACE1003850	Ó. 07
	PLACE1003858	F-PLACE1003858	0. 07
	PLACE1003864	F-PLACE1003864	0.12
10	PLACE1003900	F-PLACE1003900	0.11
	PLACE1003932	F-PLACE1003932	0. 05
	PLACE1004118	F-PLACE1004118	0. 3
	PLACE1004242	F-PLACE1004242	0. 05
15	PLACE1004256	F-PLACE1004256	0. 24
,,	PLACE1004257	F-PLACE1004257	0.06
	PLACE1004274	F-PLACE1004274,	0. 25
	PLACE1004284	F-PLACE1004284	0.13
	PLACE1004336	F-PLACE1004336	0. 27
20 .	PLACE1004384	F-PLACE1004384	0. 17
	PLACE1004425	F-PLACE1004425	0.12
	PLACE1004467	F-PLACE1004467	0. 2
	PLACE1004471	F-PLACE1004471	0.12
25	PLACE1004491	F-PLACE1004491	0.08
	PLACE1004506	F-PLACE1004506	
	PLACE1004518	F-PLACE1004518	0.14
	PLACE1004658	F-PLACE1004658	0.05
30	PLACE1004681	F-PLACE1004681	0.13
	PLACE1004693	F-PLACE1004693	0.11
	PLACE1004716	F-PLACE1004716	0.12
	PLACE1004773	F-PLACE1004773	0.17
35	PLACE1004793	F-PLACE1004793	0. 21
	PLACE1004813	F-PLACE1004813	0.11
	PLACE1004815	F-PLACE1004815	0.14
	PLACE1004827	F-PLACE1004827	0. 15
40	PLACE1004836	F-PLACE1004836	0. 25
40	PLACE1004838	F-PLACE1004838	0.16
	PLACE1004840	F-PLACE1004840	0.06
	PLACE1004900	F-PLACE1004900	0. 08
	PLACE1004913	F-PLACE1004913	0. 12
45	PLACE1004934	F-PLACE1004934	0. 17
	PLACE1004972	F-PLACE1004972	0.07
	PLACE1004979	F-PLACE1004979	0. 24
	PLACE1004985	F-PLACE1004985	0.05
50	PLACE1005052	F-PLACE1005052	0.17
	PLACE1005077	F-PLACE1005077	0.1
	PLACE1005085	F-PLACE1005085	0.13
	PLACE1005086	F-PLACE1005086	0. 13
55	PLACE1005108	F-PLACE1005108	0. 28

	PLACE1005111	F-PLACE1005111	0. 15
	PLACE1005128	F-PLACE1005128	0.12
	PLACE1005146	F-PLACE1005146	0. 07
5	PLACE1005162	F-PLACE1005162	0.13
	PLACE1005176		0.16
	PLACE1005409	F-PLACE1005409	0. 09
	PLACE1005453	F-PLACE1005453	0. 1
10	PLACE1005467	F-PLACE1005467	0. 28
70	PLACE1005471	F-PLACE1005471	0.06
	PLACE1005477	F-PLACE1005477	0. 12
	PLACE1005502	F-PLACE1005502	0. 1
	PLACE1005526	F-PLACE1005526	
15	PLACE1005528	F-PLACE1005528	0. 07
	PLACE1005574	F-PLACE1005574,	
	PLACE1005574	F-PLACE1005584	0. 24
	PLACE1005611	F-PLACE1005611	0. 18
20	PLACE1005611	F-PLACE1005639	0. 09
	PLACE1005666	F-PLACE1005666	0. 07
	PLACE1005730	F-PLACE1005730	0. 2
	PLACE1005730	F-PLACE1005802	0. 1
25	PLACE1005802 PLACE1005845	F-PLACE1005845	0. 09
	PLACE1005850	F-PLACE1005850	0. 22
	PLACE1005884	F-PLACE1005884	0. 08
	PLACE1005898	F-PLACE1005898	0. 12
	PLACE1005898	F-PLACE1005932	0. 12
30	PLACE1005932	F-PLACE1005968	0. 07
	PLACE1005908	F-PLACE1006002	0. 08
	PLACE1006002	F-PLACE1006003	0. 16
	PLACE1006017	F-PLACE1006017	0. 13
35	PLACE1006076	F-PLACE1006076	0. 11
	PLACE1006129	F-PLACE1006129	0. 25
	PLACE1006143	F-PLACE1006143	0.08
	PLACE1006164	F-PLACE1006164	0. 05
40	PLACE1006187	F-PLACE1006187	0. 2
	PLACE1006205	F-PLACE1006205	0. 08
	PLACE1006223	F-PLACE1006223	0. 05
	PLACE1006262	F-PLACE1006262	0.08
45	PLACE1006288	F-PLACE1006288	0.06
·	PLACE1006318		0. 12
	PLACE1006360		0. 07
	PLACE1006368		0. 27
50	PLACE1006371	F-PLACE1006371	0. 14
50	PLACE1006371		0. 09
	PLACE1006382 PLACE1006470		0. 14
	PLACE1006506		0. 24
	PLACE1006521		0. 08
55	LACETOUGE	, , , , , , , , , , , , , , , , , , , ,	

	PLACE1006534 F-PLACE1006534 (	0. 06
	PLACE1006540 F-PLACE1006540 (	0. 05
	PLACE1006617 F-PLACE1006617 (	0. 05
5	PLACE1006629 F-PLACE1006629 (	0. 2
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		0. 08
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10		0. 17
10	, <del></del>	0.15
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20	Y =	0.13
		0.08
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		0.08
25	PLACE1007132 F-PLACE1007132	0.07
	PLACE1007140 F-PLACE1007140	0. 16
	PLACE1007276 F-PLACE1007276	0.09
	PLACE1007286 F-PLACE1007286	0.05
3 <b>0</b>	PLACE1007478 F-PLACE1007478	0.19
	PLACE1007525 F-PLACE1007525	0.11
	PLACE1007557 F-PLACE1007557	0.13
	PLACE1007677 F-PLACE1007677	0.08
35	PLACE1007737 F-PLACE1007737	0.07
33	PLACE1007743 F-PLACE1007743	0.09
	PLACE1007807 F-PLACE1007807	0. 09
	PLACE1007829 F-PLACE1007829	0. 09
	PLACE1007852 F-PLACE1007852	0.14
40	PLACE1007866 F-PLACE1007866	0. 07
	PLACE1007877 F-PLACE1007877	0. 29
	PLACE1008045 F-PLACE1008045	0. 17
	PLACE1008080 F-PLACE1008080	0. 05
45	PLACE1008111 F-PLACE1008111	0. 11
	PLACE1008181 F-PLACE1008181	0. 08
	PLACE1008201 F-PLACE1008201	0. 12
	PLACE1008231 F-PLACE1008231	0. 08
50	PLACE1008244 F-PLACE1008244	0.06
	PLACE1008330 F-PLACE1008330	0.13
	PLACE1008331 F-PLACE1008331	0. 15
	PLACE1008369 F-PLACE1008369	0.06
55	PLACE1008392 F-PLACE1008392	0. 09
	•	

	PLACE1008405	F-PLACE1008405	0.07
	PLACE1008424	F-PLACE1008424	0.06
	PLACE1008424	F-PLACE1008426	0. 00
			0.17
5	PLACE1008437	F-PLACE1008437	
	PLACE1008455	F-PLACE1008455	0.07
	PLACE1008584	F-PLACE1008584	0.29
	PLACE1008621	F-PLACE1008621	0.07
10	PLACE1008625	F-PLACE1008625	0.12
	PLACE1008630	F-PLACE1008630	0.12
	PLACE1008643	F-PLACE1008643	0.11
	PLACE1008715	F-PLACE1008715	0.2
15	PLACE1008748	F-PLACE1008748	0.15
•	PLACE1008757	F-PLACE1008757	0.13
	PLACE1008798	F-PLACE1008798	
	PLACE1008807	F-PLACE1008807	0.12
00	PLACE1008851	F-PLACE1008851	0.22
20	PLACE1008920	F-PLACE1008920	0.1
	PLACE1008941	F-PLACE1008941	0.14
	PLACE1008947	F-PLACE1008947	0.05
	PLACE1009020	F-PLACE1009020	0. 14
25	PLACE1009039	F-PLACE1009039	0.16
	PLACE1009048	F-PLACE1009048	0.09
	PLACE1009050	F-PLACE1009050	0.09
	PLACE1009150	F-PLACE1009150	0.12
30	PLACE1009155	F-PLACE1009155	0.08
	PLACE1009172	F-PLACE1009172	0.07
	PLACE1009183	F-PLACE1009183	0.07
	PLACE1009200	F-PLACE1009200	0.08
35	PLACE1009246	F-PLACE1009246	0.08
	PLACE1009308	F-PLACE1009308	0. 23
	PLACE1009398	F-PLACE1009398	0. 26
	PLACE1009410	F-PLACE1009410	0. 15
40	PLACE1009477	F-PLACE1009477	0. 18
40	PLACE1009493	F-PLACE1009493	0.23
	PLACE1009539	F-PLACE1009539	0.11
	PLACE1009595	F-PLACE1009595	0.11
	PLACE1009613	F-PLACE1009613	0.08
45	PLACE1009621	F-PLACE1009621	0.19
	PLACE1009637	F-PLACE1009637	0. 27
	PLACE1009639	F-PLACE1009639	0. 27
	PLACE1009879	F-PLACE1009879	0. 07
50	PLACE1009888	F-PLACE1009888	0.06
	PLACE1009924	F-PLACE1009924	0.09
	PLACE1009925		0. 21
	PLACE1009935		0. 22
55	PLACE1009947	F-PLACE1009947	0.19

	PLACE1010069	F-PLACE1010069	0. 22
	PLACE1010083	F-PLACE1010083	0. 26
	PLACE1010089	F-PLACE1010089	0. 13
5	PLACE1010181	F-PLACE1010181	0. 07
	PLACE1010231	F-PLACE1010231	0. 17
	PLACE1010270	F-PLACE1010270	0. 08
	PLACE1010341	F-PLACE1010341	0. 1
10	PLACE1010562	F-PLACE1010562	0. 14
10	PLACE1010579	F-PLACE1010579	0. 15
	PLACE1010624	F-PLACE1010624	0. 13
	PLACE1010628	F-PLACE1010628	0. 16
	PLACE1010631	F-PLACE1010631	0. 23
15	PLACE1010662	F-PLACE1010662	0. 24
	PLACE1010702	F-PLACE1010702,	
	PLACE1010739	F-PLACE1010739	0. 13
	PLACE1010802	F-PLACE1010802	0. 17
20	PLACE1010891	F-PLACE1010891	0.09
	PLACE1010896	F-PLACE1010896	0. 22
	PLACE1010916	F-PLACE1010916	0. 15
	PLACE1010947	F-PLACE1010947	0.08
25	PLACE1011032	F-PLACE1011032	0. 17
	PLACE1011057	F-PLACE1011057	0. 1
	PLACE1011109	F-PLACE1011109	0.08
	PLACE1011133	F-PLACE1011133	0.08
30	PLACE1011143	F-PLACE1011143	0. 23
	PLACE1011165	F-PLACE1011165	0.05
•	PLACE1011185	F-PLACE1011185	0. 17
	PLACE1011203	F-PLACE1011203	0. 08
35	PLACE1011296	F-PLACE1011296	0. 13
-	PLACE1011340	F-PLACE1011340	0. 05
	PLACE1011375	F-PLACE1011375	0. 07
	PLACE1011419	F-PLACE1011419	0. 15
40	PLACE1011452	F-PLACE1011452	0. 13
10	PLACE1011465	F-PLACE1011465	0.14
	PLACE1011492	F-PLACE1011492	0. 26
	PLACE1011503	F-PLACE1011503	0.14
	PLACE1011520	F-PLACE1011520	0.12
45	PLACE1011567	F-PLACE1011567	0. 13
	PLACE1011576	F-PLACE1011576	0. 29
•	PLACE1011641	F-PLACE1011641	0.07
	PLACE1011643	F-PLACE1011643	0. 27
50	PLACE1011649	F-PLACE1011649	0. 17
	PLACE1011650	F-PLACE1011650	0.13
	PLACE1011719	F-PLACE1011719	0. 15
	PLACE1011729 PLACE1011749	F-PLACE1011729 F-PLACE1011749	0. 23 0. 12
55	FLAGE 1011/49	F-FLAUCIU11/49	0.12

	PLACE1011762	F-PLACE1011762	0.06
	PLACE1011778	F-PLACE1011778	0. 07
	PLACE1011874	F-PLACE1011874	0. 07
5	PLACE1011962	F-PLACE1011962	0.06
	PLACE1011964	F-PLACE1011964	0.13
	PLACE1011995	F-PLACE1011995	0.08
	PLACE2000003	F-PLACE2000003	0.09
	PLACE2000011	F-PLACE2000011	0.12
10	PLACE2000017	F-PLACE2000017	0. 22
	PLACE2000061	F-PLACE2000061	0.19
	PLACE2000103	F-PLACE2000103	0.16
	PLACE2000115	F-PLACE2000115	0.11
15	PLACE2000132	F-PLACE2000132	0. 24
	PLACE2000136	F-PLACE2000136,	0.09
	PLACE2000140	F-PLACE2000140	0. 24
	PLACE2000176	F-PLACE2000176	0. 25
20	PLACE2000187	F-PLACE2000187	0.16
	PLACE2000216	F-PLACE2000216	0. 15
	PLACE2000264	F-PLACE2000264	0.06
	PLACE2000305	F-PLACE2000305	0.05
25	PLACE2000317	F-PLACE2000317	0. 25
	PLACE2000335	F-PLACE2000335	0.09
	PLACE2000342	F-PLACE2000342	0. 07
	PLACE2000347	F-PLACE2000347	0.12
30	PLACE2000366	F-PLACE2000366	0.08
	PLACE2000379	F-PLACE2000379	0. 24
	PLACE2000394	F-PLACE2000394	0. 2
	PLACE2000398	F-PLACE2000398	0. 21
35	PLACE2000419	F-PLACE2000419	0. 05
	PLACE2000425	F-PLACE2000425	0.16
	PLACE2000435	F-PLACE2000435	0.05
	PLACE2000450	F-PLACE2000450	0.12
	PLACE2000455	F-PLACE2000455	0. 15
40	PLACE2000465	F-PLACE2000465	0. 12
	PLACE2000477	F-PLACE2000477	0.3
	PLACE3000004	F-PLACE3000004	0.3
	PLACE3000119	F-PLACE3000119	0.1
45	PLACE3000124	F-PLACE3000124	0.11
	PLACE3000136	F-PLACE3000136	0.13
	PLACE3000158	F-PLACE3000158	0.06
	PLACE3000169	F-PLACE3000169	0.13
50	PLACE3000199	F-PLACE3000199	0.06
	PLACE3000207		0.11
	PLACE3000208	F-PLACE3000208	0.07
	PLACE3000218		0.15
55	PLACE3000220	F-LY0E3000550	0.05

	PLACE3000230	F-PLACE3000230	0. 23
	PLACE3000271	F-PLACE3000271	0.17
	PLACE3000276	F-PLACE3000276	0. 2
5	PLACE3000341	F-PLACE3000341	0.08
	PLACE3000362	F-PLACE3000362	0. 19
	PLACE3000365	F-PLACE3000365	0.09
	PLACE3000373	F-PLACE3000373	0. 22
10	PLACE3000388	F-PLACE3000388	0.12
	PLACE3000399	F-PLACE3000399	0.14
	PLACE3000400	F-PLACE3000400	0.08
	PLACE3000401	F-PLACE3000401	0. 1
.5	PLACE3000402	F-PLACE3000402	0. 1
15	PLACE3000406	F-PLACE3000406	0.17
	PLACE3000475	F-PLACE3000475,	0. 07
	PLACE4000093	F-PLACE4000093	0. 21
	PLACE4000100	F-PLACE4000100	0.19
20	PLACE4000233	F-PLACE4000233	0.07
	PLACE4000247	F-PLACE4000247	0.14
	PLACE4000250	F-PLACE4000250	0.17
	PLACE4000252	F-PLACE4000252	0. 09
25	PLACE4000320	F-PLACE4000320	0. 23
	PLACE4000344	F-PLACE4000344	0. 21
	PLACE4000367	F-PLACE4000367	0. 18
	PLACE4000379	F-PLACE4000379	0.14
30	PLACE4000401	F-PLACE4000401	0. 13
	PLACE4000411	F-PLACE4000411	0. 11
	PLACE4000494	F-PLACE4000494	0. 19
	PLACE4000548	F-PLACE4000548	0. 27
35	THYR01000026	F-THYR01000026	0. 09
	THYR01000035	F-THYR01000035	0. 09
	THYR01000085	F-THYR01000085	0. 13
	THYR01000092	F-THYR01000092	0. 21
40	THYR01000111	F-THYR01000111	0. 13
	THYR01000129	F-THYR01000129	0.11
	THYR01000132	F-THYR01000132	0.1
	THYR01000156	F-THYR01000156	0. 13
	THYR01000186	F-THYR01000186	0.12
45	THYR01000187	F-THYR01000187	0.11
	THYR01000190	F-THYR01000190	0.16
	THYR01000221	F-THYR01000221	0.08
	THYR01000241	F-THYR01000241	0.08
50	THYR01000279		0.13
	THYR01000438		0.06 0.12
	THYR01000452		0. 12
	THYR01000471	F-THYR01000471	0. 11
55	THYR01000484	F-THYR01000484	V. 44

	THYR01000558	F-THYR01000558	0.09
	THYR01000596	F-THYR01000596	0.14
	THYR010C0602	F-THYR01000602	0.13
5	THYR01000625	F-THYR01000625	0. 2
	THYR01000641	F-THYR01000641	0.06
	THYR01000658	F-THYR01000658	0.12
	THYR01000699	F-THYR01000699	0.09
10	THYR01000748	F-THYR01000748	0. 3
10	THYR01000793	F-THYR01000793	0. 25
	THYR01000796	F-THYR01000796	0.11
	THYR01000805	F-THYR01000805	0. 28
	THYR01000815	F-THYR01000815	0.14
15	THYR01000843	F-THYR01000843	0.13
	THYR01000852	F-THYR01000852,	0.05
	THYR01000865	F-THYR01000865	0.14
	THYR01000895	F-THYR01000895	0.12
20	THYR01000974	F-THYR01000974	0. 07
	THYR01000975	F-THYR01000975	0.08
	THYR01001003	F-THYR01001003	0.11
	THYR01001031	F-THYR01001031	0. 1
25	THYR01001062	F-THYR01001062	0.14
	THYR01001093	F-THYR01001093	0.13
	THYR01001121	F-THYR01001121	0.14
	THYR01001133	F-THYR01001133	0. 24
30	THYR01001177	F-THYR01001177	
	THYR01001262	F-THYR01001262	0. 2
	THYR01001290	F-THYR01001290	0.06
	THYR01001321	F-THYR01001321	0.09
35	THYR01001322	F-THYR01001322	0.13
	THYR01001365	F-THYR01001365	0. 08
	THYR01001401	F-THYR01001401	0. 21
	THYR01001411	F-THYR01001411	0.06
40	THYR01001426	F-THYR01001426	0. 1
10	THYR01001434	F-THYR01001434	0.15
	THYR01001480	F-THYR01001480	0.1
	THYR01001534	F-THYR01001534	0. 26
_	THYR01001541	F-THYR01001541	0. 22
45	THYR01001559	F-THYR01001559	0.15
	THYR01001570	F-THYR01001570	0. 28
	THYR01001573	F-THYR01001573	0.05
	THYR01001595	F-THYR01001595	0.05
50	THYR01001617	F-THYR01001617	0.1
	THYR01001673	F-THYR01001673	0.07
	THYR01001706	F-THYR01001706 F-THYR01001745	0. 15 0. 12
	THYR01001745 THYR01001746	F-THYR01001745	0.12
55	101701001746	F-111KU1UU1/46	V. I

		THYR01001772	F-THYR01001772	0.08
		THYR01001895	F-THYR01001895	0.19
		THYR01001907	F-THYR01001907	0.17
5		Y79AA1000033	F-Y79AA1000033	0.18
		Y79AA1000065	F-Y79AA1000065	0. 22
		Y79AA1000346	F-Y79AA1000346	0. 2
10		Y79AA1000405	F-Y79AA1000405	0. 07
10		Y79AA1000410	F-Y79AA1000410	0.08
		Y79AA1000538	F-Y79AA1000538	0.06
		Y79AA1000802	F-Y79AA1000802	0. 28
15		Y79AA1000805	F-Y79AA1000805	0. 2
		Y79AA1000969	F-Y79AA1000969	0. 07
		Y79AA1001061	F-Y79AA1001061	0. 22
		Y79AA1001167	F-Y79AA1001167,	0. 22
20		Y79AA1001384	F-Y79AA1001384	0. 18
		Y79AA1001594	F-Y79AA1001594	0. 22
		Y79AA1001692	F-Y79AA1001692	0. 2
		Y79AA1001848	F-Y79AA1001848	0. 21
25	i	Y79AA1001875	F-Y79AA1001875	0.09
		Y79AA1002103	F-Y79AA1002103	0.11
		Y79AA1002211	F-Y79AA1002211	0. 19
30	)	Y79AA1002220	F-Y79AA1002220	0. 25
		· · ·		

# Table 11 574.clones belonging to groups (4), (5), (6), and (7), each having the maximal ATGpr1 score of not more than 0.3 and selected based on the maximal ATGpr2 score, and their maximal ATGpr2 scores

10		name of sequence	maximal ATGpr2 score
15	HEMBA1000180 HEMBA1000251 HEMBA1000280 HEMBA1000282	F-HEMBA1000180 F-HEMBA1000251 F-HEMBA1000280 F-HEMBA1000282	0. 37 0. 14 0. 33 0. 36
20	HEMBA1000333 HEMBA1000351 HEMBA1000357 HEMBA1000376	F-HEMBA1000333 F-HEMBA1000351 F-HEMBA1000357 F-HEMBA1000376 F-HFMBA1000469	0. 59 0. 36 0. 39 0. 53 0. 32
25	HEMBA1000469 HEMBA1000504 HEMBA1000519 HEMBA1000545 HEMBA1000557	F-HEMBA1000504 F-HEMBA1000519 F-HEMBA1000545 F-HEMBA1000557	0. 32 0. 34 0. 72 0. 32 0. 35 0. 34
30	HEMBA IOUUS / S	F TILMDATOOO373	J. J <del>.</del>

	HEMBA1000604	F-HEMBA1000604	0. 28
	HEMBA1000622	F-HEMBA1000622	0. 32
	HEMBA1000673	F-HEMBA1000673	0.14
5	HEMBA1000726	F-HEMBA1000726	0. 32
	HEMBA1000749	F-HEMBA1000749	0. 35
	HEMBA1000822	F-HEMBA1000822	0. 42
	HEMBA1000876	F-HEMBA1000876	0. 44
10	HEMBA1000934	F-HEMBA1000934	0.41
	HEMBA1000943	F-HEMBA1000943	0. 34
	HEMBA1000960	F-HEMBA1000960	0. 34
	HEMBA1000972	F-HEMBA1000972	0. 44
15	HEMBA1001020	F-HEMBA1001020	0. 31
	HEMBA1001024	F-HEMBA1001024	0. 22
	HEMBA1001026	F-HEMBA1001026,	
	HEMBA1001051	F-HEMBA1001051	0. 36
	HEMBA1001060	F-HEMBA1001060	0. 34
20	HEMBA1001071	F-HEMBA1001071	0. 57
	HEMBA1001208	F-HEMBA1001208	0. 57
	HEMBA1001226	F-HEMBA1001226	0. 44
	HEMBA1001319	F-HEMBA1001319	0. 58
25	HEMBA1001383	F-HEMBA1001383	0. 35
	HEMBA1001391	F-HEMBA1001391	0. 42
	HEMBA1001411	F-HEMBA1001411	0. 55
	HEMBA1001433	F-HEMBA1001433	0. 66
30 .	HEMBA1001435	F-HEMBA1001435	0.72
	HEMBA1001478	F-HEMBA1001478	0. 36
	HEMBA1001522	F-HEMBA1001522	0.31
	HEMBA1001636	F-HEMBA1001636	0. 32
35	HEMBA1001651	F-HEMBA1001651	0. 54
	HEMBA1001658	F-HEMBA1001658	0. 38
	HEMBA1001709	F-HEMBA1001709	0. 55
	HEMBA1001734	F-HEMBA1001734	0. 13
40	HEMBA1001745	F-HEMBA1001745	0. 33
	HEMBA1001784	F-HEMBA1001784	0.08
	HEMBA1001791	F-HEMBA1001791	0. 48
	HEMBA1001803	F-HEMBA1001803	0. 21
45	HEMBA1001808	F-HEMBA1001808	0. 23
	HEMBA1001835	F-HEMBA1001835	0. 67
	HEMBA1001844	F-HEMBA1001844	0. 46
	HEMBA1001888	F-HEMBA1001888	0.72
50	HEMBA1001918	F-HEMBA1001918	0. 25
<del></del>	HEMBA1001940	F-HEMBA1001940	0. 40
•	HEMBA1001962	F-HEMBA1001962	0. 63
	HEMBA1002022	F-HEMBA1002022	0. 32
	HEMBA1002185	F-HEMBA1002185	0. 33
55	HEMBA1002257	F-HEMBA1002257	0.13

	HEMBA1002348	F-HEMBA1002348	0.39
	HEMBA1002621	F-HEMBA1002621	0.47
	HEMBA1002645	F-HEMBA1002645	0.34
5	HEMBA1002661	F-HEMBA1002661	0.43
	HEMBA1002728	F-HEMBA1002728	0. 45
	HEMBA1002742	F-HEMBA1002742	0.49
	HEMBA1002921	F-HEMBA1002921	0.79
10	HEMBA1002924	F-HEMBA1002924	0.45
	HEMBA1002934	F-HEMBA1002934	0.34
	HEMBA1002968	F-HEMBA1002968	0.39
	HEMBA1003037	F-HEMBA1003037	0.43
15	HEMBA1003064	F-HEMBA1003064	0.19
	HEMBA1003071	F-HEMBA1003071	0.48
	HEMBA1003166	F-HEMBA1003166,	0.32
	HEMBA1003202	F-HEMBA1003202	0.38
20	HEMBA1003204	F-HEMBA1003204	0.34
	HEMBA1003220	F-HEMBA1003220	0.39
	HEMBA1003229	F-HEMBA1003229	0.43
	HEMBA1003276	F-HEMBA1003276	0.48
05	HEMBA1003296	F-HEMBA1003296	0.36
25	HEMBA1003373	F-HEMBA1003373	0.48
	HEMBA1003531	F-HEMBA1003531	0.46
	HEMBA1003640	F-HEMBA1003640	0.32
	HEMBA1003714	F-HEMBA1003714	0.16
30	HEMBA1003856	F-HEMBA1003856	0.47
	HEMBA1003902	F-HEMBA1003902	0.17
	HEMBA1003908	F~HEMBA1003908	0.44
	HEMBA1003926	F-HEMBA1003926	0.36
35	HEMBA1003942	F-HEMBA1003942	0. 47
	HEMBA1003987	F-HEMBA1003987	0.39
	HEMBA1004000	F-HEMBA1004000	0. 87
	HEMBA1004024	F-HEMBA1004024	0.82
40	HEMBA1004164	F-HEMBA1004164	0. 21
	HEMBA1004267	F-HEMBA1004267	0.46
	HEMBA1004323	F-HEMBA1004323	0.37
	HEMBA1004405	F-HEMBA1004405	0. 23
45	HEMBA1004433	F-HEMBA1004433	0.51
	HEMBA1004577	F-HEMBA1004577	0.51
	HEMBA1004730	F-HEMBA1004730	0.55
	HEMBA1004778	F-HEMBA1004778	0.47
50	HEMBA1004803	F-HEMBA1004803	0.36
	HEMBA1004807	F-HEMBA1004807	0.48
	HEMBA1004880	F-HEMBA1004880	0.43
	HEMBA1004900	F-HEMBA1004900	0.40
55	HEMBA1004909	F-HEMBA1004909	0.24
	HEMBA1004983	F-HEMBA1004983	0. 35

	HEMBA1005123	F-HEMBA1005123	0. 35
	HEMBA1005232	F-HEMBA1005232	0.09
	HEMBA1005241	F-HEMBA1005241	0.34
5	HEMBA1005311	F-HEMBA1005311_	0. 37
	HEMBA1005318	F-HEMBA1005318	0. 59
	HEMBA1005353	F-HEMBA1005353	0.34
	HEMBA1005374	F-HEMBA1005374	0. 38
10	HEMBA1005447	F-HEMBA1005447	0. 34
	HEMBA1005508	F-HEMBA1005508	0. 14
	HEMBA1005577	F-HEMBA1005577	0. 22
	HEMBA1005588	F-HEMBA1005588	0. 54
15	HEMBA1005593	F-HEMBA1005593	0.44
	HEMBA1005606	F-HEMBA1005606	0. 38
	HEMBA1005679	F-HEMBA1005679,	0. 87
	HEMBA1005894	F-HEMBA1005894	0. 46
20	HEMBA1005911	F-HEMBA1005911	0. 56
	HEMBA1006036	F-HEMBA1006036	0. 87
	HEMBA1006124	F-HEMBA1006124	0. 35
	HEMBA1006253	F-HEMBA1006253	0. 35
25	HEMBA1006259	F-HEMBA1006259	0. 54
	HEMBA1006364	F-HEMBA1006364	0. 55
	HEMBA1006380	F-HEMBA1006380	0. 48
	HEMBA1006426	F-HEMBA1006426	0. 54
30	HEMBA1006461	F-HEMBA1006461	0. 20
	HEMBA1006562	F-HEMBA1006562	0. 72
	HEMBA1006597	F-HEMBA1006597	0.64
	HEMBA1006639	F-HEMBA1006639	0. 55
35	HEMBA1006653	F-HEMBA1006653	0.34
33	HEMBA1006695	F-HEMBA1006695	0. 16 0. 37
	HEMBA1006696 HEMBA1006744	F-HEMBA1006696 F-HEMBA1006744	0.37
	HEMBA1006744	F-HEMBA1006744	0. 33
10	HEMBA1006949	F-HEMBA1006949	0. 78
40	HEMBA1007078	F-HEMBA1007078	0. 44
	HEMBA1007129	F-HEMBA1007129	0. 55
	HEMBA1007147	F-HEMBA1007147	0.41
45	HEMBA1007206		
45	HEMBA1007279		0. 45
	HEMBA1007327		0.71
	HEMBB1000005		0. 35
50	HEMBB1000054	F-HEMBB1000054	0.18
50	HEMBB1000055		0.87
	HEMBB1000144	F-HEMBB1000144	0.47
	HEMBB1000258	F-HEMBB1000258	0.87
	HEMBB1000318	F-HEMBB1000318	0. 34
55	HEMBB1000335	F-HEMBB1000335	0. 48

	HEMBB1000336	F-HEMBB1000336	0. 35
	HEMBB1000354	F-HEMBB1000354	0. 47
	HEMBB1000374	F-HEMBB1000374	0. 69
5	HEMBB1000402	F-HEMBB1000402	0. 54
	HEMBB1000404	F-HEMBB1000404	0. 47
	HEMBB1000480	F-HEMBB1000480	0. 35
	HEMBB1000493	F-HEMBB1000493	0. 33
10	HEMBB1000554	F-HEMB81000554	0.46
	HEMBB10C0573	F-HEMBB1000573	0.34
	HEMBB1000649	F-HEMBB1000649	0. 41
	HEMBB1000652	F-HEMBB1000652	0.46
15	HEMBB1000709	F-HEMBB1000709	0.56
	HEMB81000749	F-HEMBB1000749	0.36
	HEMBB1000790	F-HEMBB1000790,	0.58
	HEMBB1000827	F-HEMBB1000827	0.31
20	HEMBB1000831	F-HEMBB1000831	0.42
20	HEMBB1000893	F-HEMBB1000893	0.36
	HEMBB1001004	F-HEMBB1001004	0.43
	HEMBB1001008	F-HEMBB1001008	0.44
	HEMBB1001047	F-HEMBB1001047	0.47
25	HEMBB1001119	F-HEMBB1001119	0. 25
	HEMBB1001142	F-HEMBB1001142	0. 15
	HEMBB1001315	F-HEMBB1001315	0.72
	HEMBB1001317	F-HEMBB1001317	0.49
30	HEMBB1001326	F-HEMBB1001326	0.44
	HEMBB1001337	F-HEMBB1001337	0.18
	HEMBB1001367	F-HEMBB1001367	0.35
	HEMBB1001424	F-HEMBB1001424	0.34
35	HEMBB1001436	F-HEMBB1001436	0.34
	HEMBB1001458	F-HEMB81001458	0.47
	HEMBB1001535	F-HEMBB1001535	0.34
	HEMBB1001536	F-HEMBB1001536	0. 22
40	HEMBB1001565	F-HEMB81001565	0.35
	HEMBB1001747	F-HEMBB1001747	0.40
	HEMBB1001749	F-HEMBB1001749 F-HEMBB1001797	0.39 0.79
	HEMBB1001797	• • • • • • • • • • • • • • • • • • • •	0. 79
45	HEMBB1001836	F-HEMBB1001836 F-HEMBB1001863	0. 32
	HEMBB1001863	F-HEMBB1001868	0. 27
	HEMBB1001868 HEMBB1001875	F-HEMBB1001875	0. 46
	HEMBB1001911	F-HEMBB1001911	0.87
50	HEMBB1001911		0. 38
	HEMBB1001925		0. 33
	HEMBB1001944		0. 35
	HEMBB1001983		0. 32
55	HEMBB1001996	• • • • • • • • • • • • • • • • • • • •	0. 39
	,12,1001001001		

	HEMBB1001997	F-HEMBB1001997	0. 36
	HEMBB1002045	F-HEMBB1002045	0. 20
	HEMBB1002092	F-HEMBB1002092	0.31
5	HEMB81002247	F-HEMBB1002247_	0, 53
	HEMBB1002266	F-HEMBB1002266	0.34
	HEMBB1002387	F-HEMBB1002387	0. 32
	HEMBB1002425	F-HEMBB1002425	0. 31
10	HEMBB1002458	F-HEMBB1002458	0. 44
	HEMBB1002522	F-HEMBB1002522	0.56
	HEMBB1002534	F-HEMBB1002534	0.43
	HEMB81002579	F-HEMBB1002579	0. 21
15	HEMBB1002582	F-HEMBB1002582	0.48
	HEMBB1002596	F-HEMBB1002596	0. 42
	HEMBB1002617	F-HEMBB1002617,	0.47
	HEMBB1002702	F-HEMBB1002702	0.31
20	MAMMA1000043	F-MAMMA1000043	0. 45
20	MAMMA1000092	F-MAMMA1000092	0.39
	MAMMA1000129	F-MAMMA1000129	0.47
•	MAMMA1000175	F-MAMMA1000175	0. 18
	8610001VWWW	F-MAMMA1000198	0.46
25	MAMMA1000221	F-MAMMA1000221	0.43
	MAMMA1000307	F-MAMMA1000307	0.35
	MAMMA1000331	F-MAMMA1000331	0.41
	MAMMA1000356	F-MAMMA1000356	0.41
30	MAMMA1000360	F-MAMMA1000360	0. 33
	MAMMA1000402	F-MAMMA1000402	0. 35
	MAMMA1000414	F-MAMMA1000414	0. 32
	MAMMA1000444	F-MAMMA1000444	0. 13
35	MAMMA1000500	F-MAMMA1000500	0.36
	MAMMA1000522	F-MAMMA1000522	0.46
	MAMMA1000576	F-MAMMA1000576	0.58
	MAMMA1000594	F-MAMMA1000594 F-MAMMA1000597	0. 69 0. 35
40	MAMMA1000597	F-MAMMA1000720	0. 43
	MAMMA1000720	F-MAMMA1000761	0.43
	MAMMA1000761	F-MAMMA1000761	0. 15
	MAMMA1000775 MAMMA1000778	F-MAMMA1000778	0. 54
45	MAMMA1000778	F-MAMMA1000778	0. 40
	MAMMA1000798	F-MAMMA1000798	0. 40
	MAMMA1000876	F-MAMMA1000876	0. 47
	MAMMA1000870	F-MAMMA1000883	0. 41
50	MAMMA1000883		0. 45
	MAMMA1000931	F-MAMMA1000940	0. 35
	MAMMA1000941	F-MAMMA1000941	0. 35
	MAMMA1000943	F-MAMMA1000943	0. 23
55	MAMMA1000975	F-MAMMA1000975	0. 32

	MAMMA1001038	F-MAMMA1001038	0. 41
	MAMMA1001162	F-MAMMA1001162	0. 18
	MAMMA1001186	F-MAMMA1001186	0. 10
5	MAMMA1001220	F-MAMMA1001220	
	MAMMA1001256	F-MAMMA1001256	0. 45
	MAMMA1001236	F-MAMMA1001274	0. 35
		F-MAMMA1001274	0. 45
	MAMMA1001341		
10	MAMMA1001397	F-MAMMA1001397 F-MAMMA1001420	0. 64 0. 41
	MAMMA1001420	F-MAMMA1001547	
	MAMMA1001547	F-MAMMA1001590	0. 39 0. 46
	MAMMA1001590		
15	MAMMA1001670	F-MAMMA1001670	0.36
	MAMMA1001679	F-MAMMA1001679	0.46
	MAMMA1001711	F-MAMMA1001711,	
	MAMMA1001745	F-MAMMA1001745	0. 53
20	MAMMA1001760	F-MAMMA1001760	0.58
	MAMMA1001769	F-MAMMA1001769	0, 33
	MAMMA1001815	F-MAMMA1001815	0.43
	MAMMA1001820	F-MAMMA1001820	0.19
25	MAMMA1001907	F-MAMMA1001907	0.38
	MAMMA1002056	F-MAMMA1002056	0.35
	MAMMA1002078	F-MAMMA1002078	0.34
	MAMMA1002093	F-MAMMA1002093	0.55
30	MAMMA1002125	F-MAMMA1002125 F-MAMMA1002132	0.39 0.57
	MAMMA1002132 MAMMA1002145	F-MAMMA1002132	0. 78
	MAMMA1002145	F-MAMMA1002145	0.78
	MAMMA1002230	F-MAMMA10022311	0. 72
35	MAMMA1002311	F-MAMMA1002360	0. 19
33	MAMMA1002300	F-MAMMA1002411	0. 19
	MAMMA1002411	F-MAMMA1002411	0. 37
	MAMMA1002571	F-MAMMA1002571	0. 37
	MAMMA1002701	F-MAMMA1002701	0. 55
40	MAMMA1002701	F-MAMMA1002727	0. 34
	MAMMA1002728	F-MAMMA1002728	0. 55
	MAMMA1002746	F-MAMMA1002746	0. 45
	MAMMA1002764	F-MAMMA1002764	
45	MAMMA1002765	F-MAMMA1002765	0. 40
	MAMMA1002820	F-MAMMA1002820	0. 32
	MAMMA1002830	F-MAMMA1002830	0. 36
	MAMMA1002909	F-MAMMA1002909	0. 64
50	MAMMA1002941	F-MAMMA1002941	0. 42
	MAMMA1002972	F-MAMMA1002972	0. 42
	MAMMA1002973	F-MAMMA1002973	0. 35
	MAMMA1003004	F-MAMMA1003004	0. 42
55	MAMMA1003007	F-MAMMA1003007	0. 41
		, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	V. 71

	MAMMA1003039	F-MAMMA1003039	0.46
	MAMMA1003089	F-MAMMA1003089	0.58
	NT2RM4000086	F-NT2RM4000086	0.36
5	NT2RM4000265	F-NT2RM4000265	
•	NT2RM4000205	F-NT2RM4000265	
	NT2RM4000388		0. 21
	****	F-NT2RM4000414	0. 35
	NT2RM4000425	F-NT2RM4000425	0. 22
10	NT2RM4000779	F-NT2RM4000779	0. 43
	NT2RM4000855	F-NT2RM4000855	0. 52
	NT2RM4001160	F-NT2RM4001160	0. 36
	NT2RM4001313	F-NT2RM4001313	0. 38
15	NT2RM4001437	F-NT2RM4001437	0. 53
	NT2RM4001754	F-NT2RM4001754	0. 35
	NT2RM4001856	F-NT2RM4001856,	0. 27
	NT2RM4001953	F-NT2RM4001953	0. 59
20	NT2RM4001984	F-NT2RM4001984	0. 35
	NT2RM4002390	F-NT2RM4002390	0. 24
	NT2RP2000077	F-NT2RP2000077	0. 35
	NT2RP2000108	F-NT2RP2000108	0.08
25	NT2RP2000183	F-NT2RP2000183	0.43
20	NT2RP2000257	F-NT2RP2000257	0.07
	NT2RP2000289	F-NT2RP2000289	0. 35
	NT2RP2000420	F-NT2RP2000420	0. 48
	NT2RP2000678	F-NT2RP2000678	0. 55
30	NT2RP2000715	F-NT2RP2000715	0.37
	NT2RP2000842	F-NT2RP2000842	0.40
·	NT2RP2000970	F-NT2RP2000970	0.46
	NT2RP2001149	F-NT2RP2001149	0.31
35	NT2RP2001226	F-NT2RP2001226	0.44
	NT2RP2001295	F-NT2RP2001295	0.40
	NT2RP2001347	F-NT2RP2001347	0.41
	NT2RP2001467	F-NT2RP2001467	0.44
40	NT2RP2001506	F-NT2RP2001506	0.17
	NT2RP2001569	F-NT2RP2001569	0.31
	NT2RP2001663	F-NT2RP2001663	0.37
	NT2RP2001936	F-NT2RP2001936	0.33
45	NT2RP2002041	F-NT2RP2002041	0.31
	NT2RP2002047	F-NT2RP2002047	0.15
	NT2RP2002066	F-NT2RP2002066	0.11
	NT2RP2002172	F-NT2RP2002172	0.32
	NT2RP2002219	F-NT2RP2002219	0.54
50	NT2RP2002316	F-NT2RP2002316	0.39
	NT2RP2002475	F-NT2RP2002475	0.09
	NT2RP2002546	F-NT2RP2002546	0.33
	NT2RP2002591	F-NT2RP2002591	0.38
55	NT2RP2002643	F-NT2RP2002643	0. <b>5</b> 5

	NT2RP2002741	F-NT2RP2002741	0. 73
	NT2RP2002750	F-NT2RP2002750	0.48
	NT2RP2002730	F-NT2RP2002778	0.40
5	NT2RP2002778		
		F-NT2RP2002857	
	NT2RP2003000	F-NT2RP2003000	0.56
	NT2RP2003073	F-NT2RP2003073	0.36
	NT2RP2003237	F-NT2RP2003237	0.69
10	NT2RP2003394	F-NT2RP2003394	0. 34
	NT2RP2003517	F-NT2RP2003517	0. 44
	NT2RP2003668	F-NT2RP2003668	0. 46
	NT2RP2003988	F-NT2RP2003988	0. 55
15	NT2RP2004232	F-NT2RP2004232	0. 34
	NT2RP2004400	F-NT2RP2004400	0. 18
	NT2RP2004512	F-NT2RP2004512,	
	NT2RP2004523	F-NT2RP2004523	0.72
20	NT2RP2004587	F-NT2RP2004587	0.09
	NT2RP2004736	F-NT2RP2004736	0. 70
	NT2RP2004767	F-NT2RP2004767	0. 54
	NT2RP2004775	F-NT2RP2004775	0. 53
25	NT2RP2004961	F-NT2RP2004961	0.49
25	NT2RP2004962	F-NT2RP2004962	0. 87
	NT2RP2004982	F-NT2RP2004982	0. 32
	NT2RP2005289	F-NT2RP2005289	0. 24
	NT2RP2005407	F-NT2RP2005407	0.33
30	NT2RP2005531	F-NT2RP2005531	0. 20
	NT2RP2005694	F-NT2RP2005694	0.21
	NT2RP2005726	F-NT2RP2005726	0.42
	NT2RP2005942	F-NT2RP2005942	0.10
35	NT2RP2006258	F-NT2RP2006258	0. 37
	NT2RP2006261	F-NT2RP2006261	0.42
	NT2RP2006454	F-NT2RP2006454	0.46
	NT2RP2006554	F-NT2RP2006554	0.14
40	NT2RP3000055	F-NT2RP3000055	0.31
	NT2RP3000233	F-NT2RP3000233	0.54
	NT2RP3000341	F-NT2RP3000341	0.43
	NT2RP3000418	F-NT2RP3000418	0.53
45	NT2RP3000451	F-NT2RP3000451	0.46
	NT2RP3000561	F-NT2RP3000561	0. 47
	NT2RP3000582	F-NT2RP3000582	0.44
	NT2RP3001007	F-NT2RP3001007	0. 26
50	NT2RP3001281	F-NT2RP3001281	0.36
50	NT2RP3001318	F-NT2RP3001318	0.21
	NT2RP3001339	F-NT2RP3001339	0.58
	NT2RP3001340	F-NT2RP3001340	0.41
	NT2RP3001383	F-NT2RP3001383	0.36
55	NT2RP3001432	F-NT2RP3001432	0.34

	NT2RP3001580	F-NT2RP3001580	0. 45
	NT2RP3001589	F-NT2RP3001589	0.48
	NT2RP3001898	F-NT2RP3001898	0.19
5	NT2RP3002004	F-NT2RP3002004	0.36
	NT2RP3002173	F-NT2RP3002173	0.39
•	NT2RP3003133	F-NT2RP3003133	0.35
	NT2RP3003264	F-NT2RP3003264	0.13
10	NT2RP3003346	F-NT2RP3003346	0.72
•	NT2RP3003403	F-NT2RP3003403	0.45
	NT2RP3003433	F-NT2RP3003433	0. 24
	NT2RP3003576	F-NT2RP3003576	0.35
	NT2RP3003625	F-NT2RP3003625	0. 43
15	NT2RP3003665	F-NT2RP3003665	0. 32
	NT2RP3003800	F-NT2RP3003800,	
	NT2RP3003828	F-NT2RP3003828	0. 52
	NT2RP3003842	F-NT2RP3003842	0. 23
20	NT2RP3004070	F-NT2RP3004070	0. 32
	NT2RP3004470	F-NT2RP3004470	0.46
	NT2RP4000023	F-NT2RP4000023	0. 33
	NT2RP4000035	F-NT2RP4000035	0. 56
25	NT2RP4000102	F-NT2RP4000102	0. 38
	NT2RP4000167	F-NT2RP4000167	0. 42
	NT2RP4000214	F-NT2RP4000214	0. 33
	NT2RP4000214	F-NT2RP4000214	0. 37
30	NT2RP4000218	F-NT2RP4000424	0. 48
. 00	NT2RP4000424	F-NT2RP4000424	0. 40
	NT2RP4000915	F-NT2RP4000915	0. 43
	0VARC1000017	F-0VARC1000017	0. 43
	OVARC1000017	F-0VARC1000017	0. 17
35	0VARC1000085	F-0VARC1000085	0. 17
	OVARC1000083	F-0VARC1000083	0. 58
	OVARC1000092	F-0VARC1000145	0. 53
	OVARC1000145	F-0VARC1000145	0. 33
40	OVARC1000414	F-0VARC1000414	0. 40
	0VARC1000486	F-0VARC1000486	0. 13
	OVARC1000496	F-0VARC1000496	0. 33
		F-0VARC1000948	
45	OVARC1000948		
	OVARC1001011	F-0VARC1001011	0. 42 0. 14
	OVARC1001240	F-0VARC1001240	
	OVARC1001600 OVARC1001805	F-0VARC1001600	0. 37 0. 51
50		F-0VARC1001805	
	OVARC1001813	F-0VARC1001813	0.58
	OVARC1001846	F-0VARC1001846	0.59
	PLACE1000540	F-PLACE1000540	0.35
EE	PLACE1000599	F-PLACE1000599	0.37
55	PLACE1001088	F-PLACE1001088	0.36

	PLACE1001323	F-PLACE1001323	0. 21
	PLACE1001377	F-PLACE1001377	0. 41
	PLACE1001440	F-PLACE1001440	0. 45
5	PLACE1001517	F-PLACE1001517	0. 41
	PLACE1001672	F-PLACE1001672	0. 35
	PLACE1001705	F-PLACE1001705	0.11
	PLACE1001756	F-PLACE1001756	0. 67
10	PLACE1002157	F-PLACE1002157	0. 68
	PLACE1002205	F-PLACE1002205	0.55
	PLACE1002227	F-PLACE1002227	0.14
	PLACE1002259	F-PLACE1002259	0.47
15	PLACE1002319	F-PLACE1002319	0. 29
	PLACE1002399	F-PLACE1002399	0.31
	PLACE1002477	F-PLACE1002477,	0.35
	PLACE1002500	F-PLACE1002500	0. 26
22	PLACE1002583	F-PLACE1002583	0.49
20	PLACE1002604	F-PLACE1002604	0.25
	PLACE1002772	F-PLACE1002772	0.13
	PLACE1002853	F-PLACE1002853	0.46
	PLACE1002968	F-PLACE1002968	0.33
25	PLACE1003238	F-PLACE1003238	0.46
	PLACE1003420	F-PLACE1003420	0.42
	PLACE1003478	F-PLACE1003478	0.20
	PLACE1003566	F-PLACE1003566	0.35
30	PLACE1003593	F-PLACE1003593	0.42
	PLACE1003618	F-PLACE1003618	0.36
	PLACE1004274	F-PLACE1004274	0.41
	PLACE1004681	F-PLACE1004681	0. 25
35	PLACE1004716	F-PLACE1004716	0.49
	PLACE1004773	F-PLACE1004773	0. 31
	PLACE1004815	F-PLACE1004815	0. 72
	PLACE1004836	F-PLACE1004836	0.39
40	PLACE1004913	F-PLACE1004913	0.44
	PLACE1004979	F-PLACE1004979	0.44
	PLACE1005052	F-PLACE1005052	0.63
	PLACE1005086	F-PLACE1005086	0. 57
45	PLACE1005108	F-PLACE1005108	0. 17
	PLACE1005128 PLACE1005176	F-PLACE1005128 F-PLACE1005176	0. 40 0. 48
	PLACE1005176	F-PLACE1005176	0. 48
	PLACE1005487	F-PLACE1005639	0.37
50	PLACE1005859	F-PLACE1005850	0. 54
	PLACE1005830	F-PLACE1005830	0. 20
	PLACE1006932	F-PLACE1006003	0. 20
	PLACE1006017	F-PLACE1006017	0. 42
55	PLACE1006288	F-PLACE1006288	0.34
		- 1 LAGE 1000200	J. J7

	PLACE1006318	F-PLACE1006318	0. 22
	PLACE1006368	F-PLACE1006368	0.10
	PLACE1006371	F-PLACE1006371	0. 43
5	PLACE1006506	F-PLACE1006506	0. 28
	PLACE1006629	F-PLACE1006629	
	PLACE1006795	F-PLACE1006795	0.35
	PLACE1006904	F-PLACE1006904	0. 20
10	PLACE1007478	F-PLACE1007478	0.70
•	PLACE1007557	F-PLACE1007557	0.17
	PLACE1007743	F-PLACE1007743	0. 28
	PLACE1007829	F-PLACE1007829	0.16
15	PLACE1007877	F-PLACE1007877	0.16
	PLACE1008181	F-PLACE1008181	0.39
	PLACE1008330	F-PLACE1008330,	0.86
	PLACE1008584	F-PLACE1008584	0.61
	PLACE1008630	F-PLACE1008630	0.09
20	PLACE1008715	F-PLACE1008715	0.36
	PLACE1008851	F-PLACE1008851	0.39
	PLACE1008941	F-PLACE1008941	0.42
	PLACE1009039	F-PLACE1009039	0.43
25	PLACE1009048	F-PLACE1009048	0.24
	PLACE1009493	F-PLACE1009493	0.35
	PLACE1009539	F-PLACE1009539	0.31
	PLACE1009637	F-PLACE1009637	0. 58
30	PLACE1009925	F-PLACE1009925	0. 17
	PLACE1009947	F-PLACE1009947	0.42
	PLACE1010231	F-PLACE1010231	0. 36
	PLACE1010562	F-PLACE1010562	0. 63
35	PLACE1010579	F-PLACE1010579	0. 42
	PLACE1010739	F-PLACE1010739	0.49
	PLACE1010802	F-PLACE1010802	0.41
	PLACE1010896	F-PLACE1010896	0.47
40	PLACE1011032	F-PLACE1011032	0.46
	PLACE1011109	F-PLACE1011109	0. 19
	PLACE1011185		0.66
	PLACE1011452	F-PLACE1011452	0. 72
45	PLACE1011465	F-PLACE1011465	0. 51
	PLACE1011492	F-PLACE1011492	0. 19
	PLACE1011520	F-PLACE1011520	0.54
	PLACE1011567	F-PLACE1011567 F-PLACE1011643	0.51
50	PLACE1011643		0. 21 0. 55
	PLACE10117.19 PLACE1011749	F-PLACE1011719	0. 55
	PLACE1011749 PLACE2000011	F-PLACE1011749 F-PLACE2000011	0. 24
	PLACE2000017	F-PLACE2000017	0. 48
ge .	PLACE2000017	F-PLACE2000017	0. 34
55	- ENOLZU00001	I FLAGLZUUUUUI	0. 34

	PLACE2000187	F-PLACE2000187	0. 40
	PLACE2000216	F-PLACE2000216	0. 32
	PLACE2000335	F-PLACE2000335	0. 78
5	PLACE2000347	F-PLACE2000347	0. 36
	PLACE2000366	F-PLACE2000366	0. 33
	PLACE2000394	F-PLACE2000394	0. 54
	PLACE2000398	F-PLACE2000398	0. 44
10	PLACE2000425	F-PLACE2000425	0. 47
	PLACE2000450	F-PLACE2000450	0. 53
•	PLACE2000477	F-PLACE2000477	0. 36
	PLACE3000119	F-PLACE3000119	0. 40
10	PLACE3000207	F-PLACE3000207	0. 32
15	PLACE3000230	F-PLACE3000230	0. 51
	PLACE3000271	F-PLACE3000271,	0. 41
	PLACE3000373	F-PLACE3000373	0. 37
	PLACE3000399	F-PLACE3000399	0. 42
20	PLACE3000401	F-PLACE3000401	0. 37
	PLACE3000406	F-PLACE3000406	0. 46
•	PLACE4000247	F-PLACE4000247	0. 64
	PLACE4000320	F-PLACE4000320	0.36
25	PLACE4000344	F-PLACE4000344	0.31
	PLACE4000367	F-PLACE4000367	0. 32
	PLACE4000401	F-PLACE4000401	0. 54
	PLACE4000548	F-PLACE4000548	0. 26
30	THYR01000111	F-THYR01000111	0.53
	THYRO1000129	F-THYR01000129	0. 45
	THYRO1000187	F-THYR01000187	0.46
	THYRO1000279	F-THYR01000279	0.09
35	THYR01000484	F~THYR01000484	0.36
	THYRO1000596	F-THYR01000596	0.47
	THYRO1000625	F-THYR01000625	0.59
	THYRO1000793	F-THYR01000793	0.17
40	THYR01000815	F-THYR01000815	0.58
	THYRO1000865	F-THYR01000865	0.55
	THYRO1001003	F-THYR01001003	0.34
	THYRO1001031	F-THYR01001031	0.43
45	THYRO1001133	F-THYR01001133	0. 52
43	THYR01001321	F-THYR01001321	0. 37
	THYRO1001401	F-THYR01001401	0.36
•	THYRO1001426	F-THYR01001426	0.32
	THYRO1001434	F-THYR01001434	0.39
50	THYRO1001559	F-THYR01001559	0. 45
	THYRO1001570	F-THYR01001570	0. 35
	THYRO1001706	F-THYR01001706	0. 45
	THYR01001746	F-THYR01001746	0. 43
55	THYR01001772	F-THYR01001772	0. 37

THYR01001907	F-THYR01001907	0.33
Y79AA1000033	F-Y79AA1000033	0. 53
Y79AA1000346	F-Y79AA1000346	0.53
Y79AA1000410	F-Y79AA1000410	0.13
Y79AA1000805	F-Y79AA1000805	0.32
Y79AA1001692	F-Y79AA1001692	0. 45
Y79AA1002103	F-Y79AA1002103	0. 15
Y79AA1002220	F-Y79AA1002220	0.49

#### 15 EXAMPLE 13

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Full-length sequence analysis and homology search

[0192] Full-length sequence was determined for each selected cDNA clones. The nucleotide sequence determination was performed mainly by the dye-terminator method using custom synthesized DNA primers according to the primer walking procedure (custom synthesized DNA primers were used for sequencing; sequencing reaction was performed with DNA sequencing reagent supplied by PE Biosystems according to the supplier's manual; and the samples were analyzed in an automatic sequencer made by the same supplier). Sequence determination of some clones was carried out in the same manner but using a Licor DNA sequencer. Overlapping partial nucleotide sequences, which were obtained by the above-described method, were assembled together to determine a full-length nucleotide sequence. Amino acid sequences were then deduced from the determined full-length nucleotide sequences. However, amino acid sequence is not shown for a clone of which coding region was hard to be deduced or of which amino acid sequence has less than 100 amino acid residues. SEQ ID NOs corresponding to the respective clones are indicated in Table 350 and Table 351.

30 [0193] GenBank, Swiss-Prot and UniGene were searched for the determined nucleotide sequences by BLAST analysis. Matching data of cDNA clone which exhibits higher homology and of which functions are easily predicted based on the nucleotide sequences and the deduced amino acid sequences are selected from the BLAST analysis matching data with P value of 10<sup>-4</sup> or less. The matching data selected are listed herein. The results of homology search 6, 12, 13, and 14 are indicated in the last part of this specification. However, there are some clones that did not match the criteria for judgment and such matching data of BLAST analysis are not shown herein.

#### **EXAMPLE 14**

Novel full-length cDNA clone obtained from a cDNA library prepared by oligo-capping method

[0194] A cDNA clone, NT2RP4002298, was obtained from a cDNA library, NT2RP4 (see Example 1), prepared by oligo-capping method. Analysis of the entire nucleotide sequence of the clone has revealed that the clone encodes a novel protein consisting of 775 amino acids. The ATGpr1 score at the initiation codon of the amino acid sequence is 0.16, and therefore, the fullness ratio is low. However, the sequence can be full-length.

45 [0195] The full-length nucleotide sequence of NT2RP4002298 is shown in SEQ ID NO: 12370, and the deduced amino acid sequence encoded by the clone NT2RP4002298 is shown in SEQ ID NO: 12371.

#### **EXAMPLE 15**

50 Gene expression analysis with hybridization using high density DNA filter

[0196] Nylon membrane for DNA spotting was prepared according to the following procedure. E. coli was cultured in each well of a 96-well plate (in a LB medium at 37°C for 16 hours). A small aliquot of each culture was suspended in 10 μl of sterile water in a well of a 96-well plate. The plate was heated at 100°C for 10 minutes. Then the boiled samples were analyzed by PCR reaction. PCR was performed in a 20 μl solution by using TaKaRa PCR Amplification Kit (Takara) according to the supplier's protocol. Primers used for the amplification of an insert cDNA in a plasmid were a pair of sequencing primers, ME761FW (5' tacggaagtgttacttctgc 3' / SEQ ID NO: 13290) and ME1250RV (5' tgtgggaggttttttctcta 3' / SEQ ID NO: 13291), or a pair of primers, M13M4 (5' gttttcccagtcacgac 3' / SEQ ID NO: 13292)

and M13RV (5' caggaaacagctatgac 3' / SEQ ID NO: 13293). PCR reaction was performed in a thermal cycler, GeneAmp System 9600 (PE Biosystems). The cycling profile consisted of pre-heat at 95°C for 5 mimutes; 10 cycles of denaturation at 95°C for 10 seconds, and annealing/extension at 68°C for 1 minute; 20 cycles of denaturation at 98°C for 20 seconds and annealing/extension at 60°C for 3 minutes; and final extension at 72°C for 10 minutes. After the PCR reaction, the 20 µl reaction solution was loaded onto a 1 % agarose gel and fractionated by electrophoresis. DNA on the gel was stained with ethidium bromide to confirm the amplification of cDNA. When cDNAs were barely amplified by PCR, plasmids containing the corresponding insert cDNAs were prepared by the alkali-extraction method (J. Sambrook, E. F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

[0197] Preparation of DNA array was carried out by the following procedure. An Aliquot of a DNA solution was added in each well of a 384-well plate. DNA was spoited onto a nylon membrane (Boehringer) by using a 384-pin tool of Biomek 2000 Laboratory Automation Sysytem (Beckman-Coulter). Specifically, the 384-well plate containing the DNA was placed under the 384-pin tool. The independent 384 needles were simultaneously dipped into the DNA solution for DNA deposition. The needles were gently pressed onto a nylon membrane and the DNA deposited at the tips of needles was spotted onto the membrane. Denaturation of the spotted DNA and immobilization of the DNA on the nylon membrane were carried out according to usual manners (J. Sambrook, E.F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

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[0198] Hybridization probe used was radioisotope-labeled 1st strand cDNA. The 1st strand cDNA synthesis was performed by using Thermoscript<sup>(TM)</sup> RT-PCR System (GIBCO). Specifically, the 1st strand cDNA was synthesized by using 1.5  $\mu$ g mRNAs from various human tissues (Clontech), 1  $\mu$ l aliquots of 50  $\mu$ M Oligo(dT)20 and 50  $\mu$ Ci [ $\alpha$  <sup>33</sup>P] dATP according to an attached protocol. Probe purification was carried out by using ProbeQuant<sup>(TM)</sup> G-50 micro column (Amersham-Pharmacia Biotech) according to an attached protocol. In the next step, 2 units of E. coli RNasH were added to the reaction mixture. The mixture was incubated at room temperature for 10 minutes and then 100  $\mu$ g of human COT-1 DNA (GIBCO) was added thereto. The mixture was incubated at 97°C for 10 minutes and then was allowed to stand on ice to give hybridization probe.

[0199] Hybridization of the radioisotope-labeled probe to the DNA array was performed in a usual manner (J. Sambrook, E.F., Fritsh, & T. Maniatis, Molecular Cloning, A laboratory manual/2nd edition, Cold Spring Harbor Laboratory Press, 1989). The membrane was washed as follows: the nylon membrane was incubated in Washing solution 1 (2 × SSC, 1% SDS) at room temperature (about 26°C) for 20 minutes and this washing was repeated 3 times; then the membrane was washed 3 times by incubating it in Washing solution 2 (0.1  $\times$  SSC, 1% SDS) at 65°C for 20 minutes. Autoradiography was performed by using an image plate for BAS2000 (Fuji Photo Film Co., Ltd.). Specifically, the nylon membrane with probe hybridized thereon was wrapped with a piece of Saran Wrap and brought into tight contact with the image plate on the light-sensitive surface. The membrane with the image plate was placed in an imaging cassette for radioisotope and allowed to stand in dark place for 4 hours. The radioactivity recorded on the image plate was analyzed by using BAS2000 (Fuji Photo Film Co., Ltd.). The activity was subjected to electronic conversion and recorded as an image file of autoradiogram. The signal intensity of each DNA spot was analyzed by using Visage High Density Grid Analysis Systems (Genomic Solutions Inc.). The signal intensity was converted into numerical data. The data were taken by duplicated measurements. The reproducibility was assessed by comparing the signal intensities of the corresponding spots on the duplicated DNA filters that were hybridized to a single DNA probe (Figure 2). The ratio between the corresponding spots falls within a range of 2 or less in 95% of entire spots and the correlation coefficient is r=0.97. Thus the reproducibility is assumed to be satisfactory.

[0200] The detection sensitivity in gene expression analysis was estimated by examining increases in the signal intensity of probe concentration-dependent spot in hybridization using a probe complementary to the DNA spotted on the nylon membrane. DNA used was PLACE1008092 (same as DNA deposited in GenBank under an Accession No. AF107253). The DNA array with DNA of PLACE1008092 was prepared according to the above-mentioned method. The probe used was prepared as follows: mRNA was synthesized in vitro from the clone, PLACE1008092. By using this mRNA as a template, radioisotope-labeled 1st strand cDNA was synthesized in the same manner as described above, and the cDNA was used as the probe. The cDNA PLACE1008092 was inserted into pBluescript SK(-), of which T7 promoter was ligated to the 5' end of the cDNA, to give a plasmid for in vitro synthesis of the mRNA from PLACE1008092. Specifically, the PLACE1008092 insert was cut out from pME18SFL3 carrying the cDNA at a DrallI site thereof by Xhol digestion. The resulting PLACE1008092 fragment was ligated to Xhol-predigested pBluescript SK (-) by using DNA ligation kit ver.2 (Takara). The in-vitro mRNA synthesis from PLACE1008092 inserted in pBluescript SK(-) was carried out by using Ampliscribe(TM) T7 high yield transcription kit (Epicentre technologies). Hybridization and the analysis of signal intensity of each DNA spot were conducted by the same methods as described above. When the probe concentration is 1  $\times$  10<sup>7</sup>  $\mu$ g/ml or less, there was no increase of signal intensity proportional to the probe concentration. Therefore it was assumed to be difficult to compare the signals with one another in the concentration range. Thus the spots with the intensity of 40 or less were indiscriminately taken as low-level signals (Figure 3). Within a concentration of the probe ranging from 1  $\times$  10<sup>7</sup>  $\mu$ g/ml to 0.1  $\mu$ g/ml, the signal was found to increase in a probe

concentration-dependent manner. The detection limit was then assumed to be 1:100,000 in a ratio of mRNA expression level in a sample.

[0201] Tables 12-168 (also containing clones with no description in Examples) show the expression of each cDNA in human normal tissues (heart, lung, pituitary gland, thymus, brain, kidney, liver and spleen). The expression levels are indicated by numerical values of 0-10,000. Genes that were expressed in at least a single tissue are indicated below by the corresponding clone names:

	HEMBA1000012,	HEMBA1000020,	HEMBA1000030,	HEMBA1000042,	HEMBA1000046,
10	HEMBA1000076,	HEMBA1000111,	HEMBA1000129,	HEMBA1000150,	HEMBA1000156,
	HEMBA1000158,	HEMBA1000168,	HEMBA1000185,	HEMBA1000216,	HEMBA1000227,
	HEMBA1000231,	HEMBA1000243,	HEMBA1000244,	HEMBA1000280,	HEMBA1000282,
	HEMBA1000288,	HEMBA1000303,	HEMBA1000304,	HEMBA1000327,	HEMBA1000338,
15	HEMBA1000351,	HEMBA1000355,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366,
	HEMBA1000369,	HEMBA1000376,	HEMBA1000387,	HEMBA1000392,	HEMBA1000396,
	HEMBA1000411,	HEMBA1000422,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000469,	HEMBA1000488,	HEMBA1000491,	HEMBA1000501,	HEMBA1000505,
20	HEMBA1000508,	HEMBA1000519,	HEMBA1000523,	HEMBA1000534,	HEMBA1000540,
	HEMBA1000542,	HEMBA1000545,	HEMBA1000557,	HEMBA1000561,	HEMBA1000568,
	HEMBA1000569,	HEMBA1000575,	HEMBA1000588,	HEMBA1000591,	HEMBA1000604,

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	-HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000657,	HEMBA1000673,
	HEMBA1000682,	HEMBA1000702,	HEMBA1000705,	HEMBA1000726,	HEMBA1000749,
	HEMBA1000752,	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,
5	HEMBA1000843,	HEMBA1000852,	HEMBA1000870,	HEMBA1000872,	HEMBA1000876,
	HEMBA1000908,	HEMBA1000910,	HEMBA1000918,	HEMBA1000919,	HEMBA1000934,
	HEMBA1000942,	HEMBA1000960,	HEMBA1000971,	HEMBA1000972,	HEMBA1000975,
	HEMBA1000986,	HEMBA1000991,	HEMBA1001007,	HEMBA1001008,	HEMBA1001017,
10	HEMBA1001020.	HEMBA1001051, HEM			•
	HEMBA1001080,	HEMBA1001085,	HEMBA1001088,	HEMBA1001109,	HEMBA1001121,
	HEMBA1001123,	HEMBA1001133,	HEMBA1001137,	HEMBA1001140,	HEMBA1001172,
	HEMBA1001197,	HEMBA1001208,	HEMBA1001213,	HEMBA1001226,	HEMBA1001235,
15	HEMBA1001265,	HEMBA1001281,	HEMBA1001286,	HEMBA1001289,	HEMBA1001294,
,,,	HEMBA1001299,	HEMBA1001302,	HEMBA1001303,	HEMBA1001310,	HEMBA1001326,
	HEMBA1001330,	HEMBA1001351,,	HEMBA1001375,	HEMBA1001377,	HEMBA1001387,
	HEMBA1001388,	HEMBA1001398,	HEMBA1001405,	HEMBA1001413,	HEMBA1001415,
	HEMBA1001432,	HEMBA1001433,	HEMBA1001435,	HEMBA1001446,	HEMBA1001450,
20	HEMBA1001454,	HEMBA1001463,	HEMBA1001476,	HEMBA1001497,	HEMBA1001510-
	HEMBA1001515,	HEMBA1001517,	HEMBA1001526,	HEMBA1001533,	HEMBA1001557,
	HEMBA1001566,	HEMBA1001569,	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,
	HEMBA1001589,	HEMBA1001595,	HEMBA1001608,	HEMBA1001620,	HEMBA1001636;
25	HEMBA1001640.	HEMBA1001647,	HEMBA1001651,	HEMBA1001655,	HEMBA1001661,
	HEMBA1001678,	HEMBA1001709,	HEMBA1001712.	HEMBA1001714,	HEMBA1001718,
	HEMBA1001731.	HEMBA1001734,	HEMBA1001745,	HEMBA1001761,	HEMBA1001781,
	HEMBA1001784,	HEMBA1001791,	HEMBA1001804,	HEMBA1001808,	HEMBA1001809,
30	HEMBA1001815,	HEMBA1001822,	HEMBA1001824,	HEMBA1001844,	HEMBA1001847,
	HEMBA1001864,	HEMBA1001866,	HEMBA1001869,	HEMBA1001888,	HEMBA1001896,
	HEMBA1001910,	HEMBA1001912,	HEMBA1001913,	HEMBA1001939,	HEMBA1001940,
	HEMBA1001964,	HEMBA1001967, HE	MBA1001987, HEMB	A1001991, HEMBA1	002003,
35	HEMBA1002008,	HEMBA1002018,	HEMBA1002035,	HEMBA1002049,	HEMBA1002092,
	HEMBA1002100,	HEMBA1002102,	HEMBA1002113,	HEMBA1002119,	HEMBA1002125,
	HEMBA1002144,	HEMBA1002150,	HEMBA1002151,	HEMBA1002160,	HEMBA1002161,
	HEMBA1002162,	HEMBA1002166,	HEMBA1002177,	HEMBA1002185,	HEMBA1002189,
40	HEMBA1002191,	HEMBA1002226,	HEMBA1002229,	HEMBA1002237,	HEMBA1002241,
	HEMBA1002265,	HEMBA1002267,	HEMBA1002270,	HEMBA1002328,	HEMBA1002337,
	HEMBA1002341,	HEMBA1002363,	HEMBA1002381,	HEMBA1002389,	HEMBA1002417,
	HEMBA1002419,	HEMBA1002439,	HEMBA1002458,	HEMBA1002462,	HEMBA1002469,
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	HEMBA1002508,	HEMBA1002513,	HEMBA1002515,	HEMBA1002538,	HEMBA1002542,
	HEMBA1002552,	HEMBA1002558,	HEMBA1002561,	HEMBA1002569,	HEMBA1002590,
	HEMBA1002592,	HEMBA1002609,	HEMBA1002624,	HEMBA1002629,	HEMBA1002645,
50	HEMBA1002659,	HEMBA1002661,	HEMBA1002678,	HEMBA1002679,	HEMBA1002696,
50	HEMBA1002703,	HEMBA1002712,	HEMBA1002728,	HEMBA1002730,	HEMBA1002746,
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5 <i>5</i>	HEMBA1002921,	HEMBA1002924,	HEMBA1002934,	HEMBA1002935,	HEMBA1002944,

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	HEMBA1003046,	HEMBA1003067,	HEMBA1003071,	HEMBA1003078,	HEMBA1003079,
5	HEMBA1003083,	HEMBA1003086,	HEMBA1003098,	HEMBA1003129,	HEMBA1003133,
	HEMBA1003136,	HEMBA1003142,	HEMBA1003148,	HEMBA1003166,	HEMBA1003175,
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	HEMBA1003538,	HEMBA1003556,	HEMBA1003571,	HEMBA1003581,	HEMBA1003591,
	HEMBA1003595,	HEMBA1003597,	HEMBA1003598,	HEMBA1003615,	HEMBA1003617,
	HEMBA1003621,	HEMBA1003637,	HEMBA1003640,	HEMBA1003645,	HEMBA1003656,
20	HEMBA1003667,	HEMBA1003680,	HEMBA1003690,	HEMBA1003692,	HEMBA1003711,
	HEMBA1003714,	HEMBA1003715,	HEMBA1003720,	HEMBA1003725,	HEMBA1003729,
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	HEMBA1003803,	HEMBA1003804,	HEMBA1003805,	HEMBA1003827,	HEMBA1003836,
05	HEMBA1003838,	HEMBA1003864,	HEMBA1003879,	HEMBA1003880,	HEMBA1003885,
25	HEMBA1003893,	HEMBA1003902,	HEMBA1003926,	HEMBA1003937,	HEMBA1003942,
	HEMBA1003958,	немва1003978, нем	BA1003987, HEMB	A1003989, HEMBA1	004011,
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	HEMBA1004200,	HEMBA1004203,	HEMBA1004225,	HEMBA1004227,	HEMBA1004238,
	HEMBA1004246,	HEMBA1004267,	HEMBA1004274,	HEMBA1004275,	HEMBA1004289,
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	HEMBA1004460,	HEMBA1004461,	HEMBA1004479,	HEMBA1004499,	HEMBA1004506,
	HEMBA1004507,	HEMBA1004509,	HEMBA1004534,	HEMBA1004538,	HEMBA1004542,
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	HEMBA1004670,	HEMBA1004672,	HEMBA1004697,	HEMBA1004705,	HEMBA1004709,
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	HEMBA1004909,	HEMBA1004918,	HEMBA1004923,	HEMBA1004930,	HEMBA1004933,
50	HEMBA1004934,	HEMBA1004944,	HEMBA1004954,	HEMBA1004960,	HEMBA1004972,
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	HEMBA1005113,	HEMBA1005202,	HEMBA1005206,	HEMBA1005223,	HEMBA1005152,
55	HEMBA1005244,	HEMBA1005251,	HEMBA1005252,	HEMBA1005275,	
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HEMBA1005353, HEMBA1005359, HEMBA1005367, HEMBA1005374, HEMBA1005382, HEMBA1005403, HEMBA1005408, HEMBA1005417, HEMBA1005405, HEMBA1005405, HEMBA1005405, HEMBA1005474, HEMBA1005475, HEMBA1005510, HEMBA1005511, HEMBA1005513, HEMBA1005518, HEMBA1005518, HEMBA1005518, HEMBA1005520, HEMBA1005520, HEMBA1005527, HEMBA1005528, HEMBA1005528, HEMBA1005529, HEMBA100531, HEMBA1005529, HEMBA1005529, HEMBA100531, HEMBA100531, HEMBA1005329, HEMBA1005331, HEM		HEMBA1005304,	HEMBA10053-11,	HEMBA1005315,	HEMBA1005331,	HEMBA1005338,
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HEMBA1005443,				HEMBA1005408,	HEMBA1005411,	HEMBA1005423,
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[0202] Genes that were expressed in all the tissues tested are indicated below by the corresponding clone names:

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35	Y79AA1002482.				

[0203] Genes that were expressed at low levels in any of the tissues tested are indicated below by the corresponding clone names:

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	PLACE1009099,	PLACE1009110,	PLACE1009113,	PLACE1009130,	PLACE1009150,
	PLACE1009158,	PLACE1009172, PLA	ACE1009186, PLAC	CE1009190, PLACE1	009230,
35	PLACE1009298,	PLACE1009319,	PLACE1009328,	PLACE1009335,	PLACE1009338,
	PLACE1009375,	PLACE1009398,	PLACE1009404,	PLACE1009410,	PLACE1009434,
	PLACE1009443,	PLACE1009468,	PLACE1009476,	PLACE1009477,	PLACE1009493,
	PLACE1009539,	PLACE1009542,	PLACE1009571,	PLACE1009581,	PLACE1009596,
40	PLACE1009613,	PLACE1009622,	PLACE1009637,	PLACE1009639,	PLACE1009659,
	PLACE1009665,	PLACE1009708,	PLACE1009721,	PLACE1009763,	PLACE1009794,
•	PLACE1009798,	PLACE1009845,	PLACE1009886,	PLACE1009888,	PLACE1009908,
	PLACE1009921,	PLACE1009924,	PLACE1009925,	PLACE1009935,	PLACE1009971,
45	PLACE1009992,	PLACE1009995,	PLACE1009997,	PLACE1010031,	PLACE1010053,
	PLACE1010069,	PLACE1010083,	PLACE1010089,	PLACE1010096,	PLACE1010102,
	PLACE1010105,	PLACE1010106,	PLACE1010134,	PLACE1010148,	PLACE1010152,
	PLACE1010181,	PLACE1010194,	PLACE1010202,	PLACE1010231,	PLACE1010261,
50	PLACE1010270,	PLACE1010274,	PLACE1010324,	PLACE1010329,	PLACE1010341,
	PLACE1010364,	PLACE1010401,	PLACE1010481,	PLACE1010491,	PLACE1010492,
	PLACE1010547, PLACE1010622,	PLACE1010562,	PLACE1010579, PLACE1010628,	PLACE1010599, PLACE1010630,	PLACE1010616,
	PLACE1010622,	PLACE1010624, PLACE1010714,	PLACE1010628, PLACE1010739,	PLACE1010630, PLACE1010743,	PLACE1010631, PLACE1010800,
55	PLACE1010802,	PLACE1010714, PLACE1010811,	PLACE1010759, PLACE1010857,		PLACE1010800, PLACE1010877,
	. MIODIUIU002,	LPUCETAINOTT	r PUCETA10091'	PLACE1010870,	TEMORIUIU8//,

	PLACE1010891,	PLACE1010896,	PLACE1010916,	PLACE1010917,	PLACE1010960,
	PLACE1010965,	PLACE1011026, PLA	ACE1011057, PLAC	E1011143, PLACE1	011160,
5	PLACE1011165,	PLACE1011203,	PLACE1011214,	PLACE1011221,	PLACE1011229,
	PLACE1011263,	PLACE1011273,	PLACE1011310,	PLACE1011325,	PLACE1011332,
	PLACE1011375,	PLACE1011399,	PLACE1011419,	PLACE1011433,	PLACE1011465,
	PLACE1011503,	PLACE1011520,	PLACE1011563,	PLACE1011635,	PLACE1011641,
10	PLACE1011643,	PLACE1011664,	PLACE1011675,	PLACE1011729,	PLACE1011762,
	PLACE1011778,	PLACE1011858,	PLACE1011874,	PLACE1011875,	PLACE1011896,
	PLACE1011964,	PLACE1011982,	PLACE1012031,	PLACE2000006,	PLACE2000014,
	PLACE2000015,	PLACE2000021,	PLACE2000033,	PLACE2000061,	PLACE2000072,
15	PLACE2000097,	PLACE2000115,	PLACE2000136,	PLACE2000164,	PLACE2000223,
15	PLACE2000371,	PLACE2000379,	PLACE2000477,	PLACE3000059,	PLACE3000121,
	PLACE3000160,	PLACE3000199,	PLACE3000218,	PLACE3000244,	PLACE3000310,
	PLACE3000320,	PLACE3000339,	PLACE3000350,	PLACE3000373,	PLACE4000089,
20	PLACE4000093,	PLACE4000147,	PLACE4000252,	PLACE4000270,	PLACE4000367,
20	PLACE4000392,	PLACE4000401,	PLACE4000558,	PLACE4000590,	PLACE4000654,
	PLACE4000670,	SKNMC1000011,	SKNMC1000013,	SKNMC1000046,	SKNMC1000050,
	SKNMC1000091,	THYR01000035,	THYR01000111,	THYR01000121,	THYR01000124,
25	THYR01000129,	THYR01000156,	THYRO1000199,	THYR01000242,	THYR01000270,
23	THYR01000279,	THYR01000288,	THYR01000327,	THYR01000381,	THYR01000438,
	THYR01000488,	THYR01000501,	THYR01000502,	THYR01000505,	THYR01000570,
	THYR01000596,	THYR01000605, TH	YRO1000641, THYR	01000662, THYR01	000734,
30	THYR01000777,	THYR01000783,	THYR01000793,	THYR01000796,	THYR01000805,
	THYR01000829,	THYR01000852,	THYR01000895,	THYR01000934,	THYR01000952,
	THYR01001033,	THYR01001100,	THYR01001134,	THYR01001142,	THYR01001204,
	THYR01001262,	THYR01001271,	THYR01001290,	THYR01001347,	THYR01001559,
35	THYR01001661,	THYR01001721,	THYR01001745,	Y79AA1000328,	Y79AA1000420,
	Y79AA1000705,	Y79AA1000734,	Y79AA1000748,	Y79AA1000752,	Y79AA1000774,
	Y79AA1000784,	Y79AA1000802,	Y79AA1000824,	Y79AA1000827,	Y79AA1000976,
	Y79AA1001078,	Y79AA1001281,	Y79AA1001312,	Y79AA1001493,	Y79AA1001541,
40	Y79AA1001581,	Y79AA1001585,	Y79AA1001696,	Y79AA1001781,	Y79AA1001848,
	Y79AA1001874,	Y79AA1001923,	Y79AA1002027,	Y79AA1002115,	Y79AA1002139,
	Y79AA1002208,	Y79AA1002209,	Y79AA1002210,	Y79AA1002220,	Y79AA1002298,
	Y79AA1002307,	Y79AA1002311,	Y79AA1002407,	Y79AA1002416,	Y79AA1002431,
45	Y79AA1002487.		•		

[0204] Genes exhibiting characteristic features in the expression thereof were selected by statistical analysis of these data. Two examples are shown below to describe the selection of genes of which expression is varied greatly among tissues. The β' actin gene is used frequently as a control in gene expression analysis. Genes of which expression is varied greatly among tissues as compared that of the β-actin gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of β-actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_a^2$ . Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_b^2$ . By taking variance ratio F as  $F=S_b^2/S_a^2$ , genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below by the corresponding clone names:

HEMBA1002113, HEMBA1005296, HEMBA1007121, HEMBB1000637, HEMBB1000915, MAMMA1000597, MAMMA1000605, MAMMA1000962, MAMMA1001139, MAMMA1001198, MAMMA1002858, NT2RM2001896,

NT2RP2002710, NT2RP2004339, NT2RP2004538, NT2RP3000348, NT2RP3003121, PLACE3000009,

PLACE3000254, THYRO1000569, Y79AA1000131.

[0205] Gene of OVARC1000037 {heterogeneous nuclear ribonucleoprotein (hnRNP)} which expression is varied little. Genes of which expression is varied greatly among tissues as compared that of the OVARC1000037 gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of  $\beta$ -actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_a^2$ . Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_b^2$ . By taking variance ratio F as  $F = S_b^2 / S_a^2$ , genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below,by the corresponding clone names:

10	HEMBA1000304,	NT2RM2001716,	NT2RM2001840,	HEMBA1001051,	HEMBA1001109,
	OVARC1001731,	HEMBA1000726,	HEMBA1001286,	HEMBA1000387,	HEMBA1000519,
	NT2RM2001896.	HEMBA1000042,	HEMBA1001085,	HEMBA1001330,	0VARC1000576_
	HEMBA1000575,	NT2RM2000599,	NT2RM2000714,	HEMBA1000469,	NT2RM4000366,
15	HEMBA1001377,	HEMBA1000769,	HEMBA1000338,	NT2RM2000795,	HEMBA1001299,
	HEMBA1000508,	HEMBA1000150,	HEMBA1000774,	HEMBA1001226,	HEMBA1000960;
	NT2RM4000795,	HEMBA1002162,	NT2RM4001876,	NT2RM4002482,	HEMBA1001678,
	HEMBA1002113,	NT2RM4002383,	HEMBA1002229,	HEMBA1002818,	HEMBA1001454,
20	NT2RM4000764,	HEMBA1001510,	HEMBA1001714,	HEMBA1002150,	NT2RM4002044,
	HEMBA1002728,	NT2RM4002189,	HEMBA1001991,	HEMBA1002166,	NT2RM4002499,
	NT2RM4001140,	NT2RM4002504,	HEMBA1002590,	HEMBA1001435,	PLACE1000706,
	HEMBA1002160,	HEMBA1001824,	HEMBA1001463,	HEMBA1001533,	HEMBA1001570,
25	PLACE1001036,	HEMBA1001651,	HEMBA1002381,	HEMBA1002934,	HEMBA1003370,
	HEMBA1003021,	HEMBA1003166,	NT2RP1000738,	NT2RP2000040,	HEMBA1004164,
	HEMBA1003836,	HEMBA1004267,	NT2RP2000845,	HEMBA1003041,	HEMBA1003571,
	HEMBA1003758,	NT2RP2000108,	HEMBA1003838,	NT2RP1000357,	HEMBA1003376,
30	PLACE1003528,	HEMBA1003528,	NT2RP1001475,	HEMBA1004049,	HEMBA1003212,
	HEMBA1003667,	PLACE1004149,	HEMBA1003926,	HEMBA1004306,	HEMBA1004024,
	NT2RP1000363,	HEMBA1003033,	HEMBA1004335,	HEMBA1003348,	HEMBA1003034,
0.5	NT2RP2001081,	HEMBA1004056, HE	MBA1003314, HEME	BA1003827, HEMBA1	003893,
35	NT2RP2001036,	NT2RP2001168,	NT2RP2001328,	HEMBA1005035,	NT2RP2001569,
	NT2RP2002439,	HEMBA1005511,	HEMBA1005999,	NT2RP2002862,	NT2RP2002979,
	NT2RP2001394,	HEMBA1004753,	NT2RP2002621,	HEMBA1005853,	HEMBA1005443,
40	NT2RP2002980,	NT2RP2001347,	HEMBA1005241,	NT2RP2002750,	NT2RP2003533,
40	HEMBA1005634,	NT2RP2003034,	HEMBA1006138,	NT2RP2003117,	NT2RP2001366,
	HEMBA1005079,	NT2RP2003293,	NT2RP2002710,	HEMBA1005911,	NT2RP2002752,
	HEMBA1006036,	NT2RP2002987,	HEMBA1006100,	HEMBA1004460,	HEMBA1004538,
45	NT2RP2001943,	NT2RP2002033,	HEMBA1005296,	HEMBA1005829,	HEMBA1005520,
45	HEMBA1005123,	HEMBA1005552,	HEMBA1004930,	NT2RP2001312,	HEMBA1005304,
	HEMBA1005834,	HEMBA1005990,	HEMBA1005526,	NT2RP2003073,	HEMBA1005331,
	HEMBA1006744,	HEMBA1006780,	NT2RP2004339,	HEMBB1000173,	HEMBA1007113,
50	NT2RP2005908,	HEMBB1000376,	HEMBB1000024,	HEMBB1000510,	NT2RP2004580,
50					

	NT2RP2003912,	HEMBA1006381,	NT2RP2004270,	HEMBA1006993,	NT2RP2005325,
	HEMBB1000218,	HEMBA1007121,	HEMBB1000434,	NT2RP2006166,	PLACE1010310,
	HEMBB1000589,	NT2RP2003986,	HEMBA1006419,	HEMBB1000141,	HEMBB1000030,
5	HEMBB1000099,	HEMBA1006624,	NT2RP2003988,	NT2RP2005753,	HEMBA1007147,
	HEMBB1000354,	NT2RP2006023,	HEMBB1000441,	HEMBB1000523,	HEMBB1000059,
	HEMBA1006631,	NT2RP2004538,	HEMBA1006695,	HEMBB1000490,	NT2RP2006184,
	HEMBB1000573,	NT2RP2004675,	HEMBA1007078,	NT2RP2005457,	NT2RP2005491,
10	HEMBB1000491,	HEMBB1000575, PLA		P2005773, NT2RP2	005354,
	HEMBB1000337,	HEMBB1000008,	NT2RP2005701,	NT2RP2003825,	HEMBB1000215,
	HEMBB1000258,	NT2RP2005496,	HEMBB1000374,	NT2RP2005581,	HEMBB1000018,
	HEMBB1000493,	HEMBB1000554,	HEMBB1000671,	NT2RP3000348,	HEMBB1001267,
15	HEMBB1002092,	HEMBB1001749,	PLACE3000304,	HEMBB1001834,	HEMBB1001142,
13	HEMBB1001177,	HEMBB1000840,	HEMBB1001436,	HEMBB1001921,	НЕМВВ1002005,
	NT2RP3001398,	HEMBB1002069,	HEMBB1002094,	HEMBB1001836,	HEMBB1001114,
	NT2RP3000628,	HEMBB1001367,	HEMBB1001588,	HEMBB1002249,	HEMBB1000684,
	NT2RP2006571,	HEMBB1001802,	HEMBB1000915,	HEMBB1000996,	NT2RP3001274,
20	HEMBB1001564,	NT2RP3000968,	PLACE3000156,	HEMBB1002045,	NT2RP3000341,
	HEMBB1000917,	PLACE1011090,	NT2RP3000742,	PLACE2000435,	HEMBB1001944,
	PLACE3000254,	HEMBB1002218,	NT2RP2006436,	NT2RP3000320,	HEMBB1001126,
	HEMBB1000749,	NT2RP3000644,	HEMBB1001527,	NT2RP3000850,	HEMBB1001967,
25	PLACE3000401,	HEMBB1000637,	HEMBB1001234,	NT2RP3000080,	PLACE2000411,
	PLACE3000009,	PLACE3000070,	HEMBB1002134,	PLACE3000475,	PLACE4000192,
	HEMBB1002520,	HEMBB1002545,	MAMMA1000155,	MAMMA1000307,	THYR01000787,
	MAMMA1000348,	MAMMA1000372,	NT2RP3003121,	MAMMA1000851,	MAMMA1000501,
30	THYRO1000569,	MAMMA1000576,	MAMMA1000605,	HEMBB1002442,	HEMBB1002596,
	MAMMA1000198,	MAMMA1000714,	MAMMA1000356,	MAMMA1000760,	MAMMA1000416,
	MAMMA1000931,	MAMMA1000962,	HEMBB1002387,	NT2RP3001712,	HEMBB1002556,
	HEMBB1002617,	MAMMA1000171, PL	ACE4000612, MAMM	MAMMA1 (MAMMA)	.000421,
35	MAMMA1000444,	HEMBB1002358,	HEMBB1002453,	HEMBB1002664,	HEMBB1002699,
	MAMMA1000720,	MAMMA1000360,	MAMMA1000744,	MAMMA1000802,	MAMMA1000839,
	MAMMA1000478,	MAMMA1000877,	MAMMA1000585,	MAMMA1000941,	NT2RP3002909,
	MAMMA1000968,	HEMBB1002457,	HEMBB1002495,	MAMMA1000057,	MAMMA1000696,
40	МАММА 1000942,	MAMMA1000302,	MAMMA1000943,	MAMMA1000998,	NT2RP3001752,
40	NT2RP3003032,	MAMMA1000429,	MAMMA1000565,	MAMMA1000594,	MAMMA1000625,
	NT2RP3002146,	MAMMA1000084,	MAMMA1000257,	NT2RP3003157,	MAMMA1000431,
	MAMMA1000597,	MAMMA1001078,	MAMMA1001126,	THYR01001537,	NT2RP3004093,
	MAMMA1001202,	NT2RP3003842,	Y79AA1000342,	MAMMA1002311,	MAMMA1001969,
45	Y79AA1000560,	MAMMA1002359,	MAMMA1002056,	MAMMA1002413,	MAMMA1002454,
	MAMMA1002612,	MAMMA1001411,	MAMMA1001035,	MAMMA1001080,	MAMMA1001133,
	NT2RP3004095,	MAMMA1001161,	MAMMA1001203,	MAMMA1001663,	NT2RP3003819,
	MAMMA1001745,	MAMMA1001790,	NT2RP3004470,	NT2RP3004503,	MAMMA1002293,
50	NT2RP4000321,	MAMMA1001880,	Y79AA1000410,	MAMMA1001970,	MAMMA1002058,
	NT2RP4000515,	MAMMA1002215,	MAMMA1002617,	NT2RP3003576,	MAMMA1001139,
	MAMMA1001576,	Y79AA1000131,	MAMMA1001330,	MAMMA1001769,	MAMMA1001383,
	MAMMA1002032,	MAMMA1002125,	MAMMA1002174,	NT2RP4000614,	NT2RP4000243,
55	NT2RP3004110,	MAMMA1001271,	MAMMA1001296,	MAMMA1001992,	MAMMA1002033,

	MAMMA1002428,	MAMMA1002590,	MAMMA1001186,	MAMMA1002267,	MAMMA1002322,
	MAMMA1001956,	MAMMA1002155, NT	2RP4000210, MAMMA	A1002622, NT2RP3	004125,
_	MAMMA1001220,	MAMMA1001683,	NT2RP3004348,	Y79AA1000214,	Y79AA1000833,
5	NT2RP4000212,	MAMMA1002230,	MAMMA1001452,	MAMMA1001620,	MAMMA1001256,
	MAMMA1001760,	NT2RP3004349,	MAMMA1001783,	MAMMA1001907,	MAMMA1002009,
	MAMMA1002545,	NT2RP4000214,	NT2RP4000728,	MAMMA1001465,	MAMMA1001154,
	MAMMA1001198,	MAMMA1001343,	MAMMA1002310,	NT2RP4000035,	NT2RP4000833,
10	MAMMA1003150,	MAMMA1002886,	NT2RP4001938,	NT2RM2000260,	MAMMA1002629,
	MAMMA1002973,	MAMMA1002721,	MAMMA1002909,	NT2RP4001100,	NT2RM1000857,
	NT2RP4000878,	MAMMA1002844,	NT2RM1000039,	NT2RP4001174,	MAMMA1002665,
	MAMMA1003047,	NT2RM1000086,	NT2RM1000260,	NT2RM1000355,	MAMMA1002701,
15	NT2RP4000918,	MAMMA1002830,	MAMMA1002970,	NT2RP4001677,	NT2RM2000422,
	MAMMA1003004,	MAMMA1002673,	MAMMA1003031,	MAMMA1002764,	MAMMA1002858,
	NT2RP4001679,	NT2RP4002888,	MAMMA1002711,	NT2RP4001276,	NT2RM1000018,
	NT2RP4001568,	NT2RM1000883.			

[0206] Thus, characteristic features in the expression of a gene are illustrated by comparing and \_ statistically analyzing the expression of many genes.

#### 25 Analysis of disease-associated genes

[0207] Non-enzymic protein glycation reaction is believed to be a cause of a variety of chronic diabetic complications. Accordingly, genes of which expression is elevated or decreased in a glycated protein-specific manner in the endothelial cells are associated with diabetic complications caused by glycated proteins. Vascular endothelial cells are affected with glycated proteins present in blood. Reaction products of non-enzymic protein glycation include amadori compound (glycated protein) as a mildly glycated protein and advanced glycation endproduct as a heavily glycated protein. Hence, a survey was carried out for genes of which expression levels are varied depending on the presence of these glycated proteins in endothelial cells. The mRNAs were extracted from endothelial cells that were cultured in the presence or absence of glycated protein. The mRNAs were converted into radiolabeled first strand cDNAs for preparing probes. The probes were hybridized to the above-mentioned DNA array. Signal of each DNA spot was detected by BAS2000 and analyzed by ArrayGauge (Fuji Photo Film Co., Ltd.).

[0208] Advanced glycation endproduct of bovine serum albumin was prepared as follows: bovine serum albumin (BSA; Sigma) was incubated in a phosphate buffer solution containing 50 mM glucose at 37°C for 8 weeks; and the resulting brownish BSA was dialyzed against a phosphate buffer solution.

[0209] Human normal pulmonary arterial endothelial cells (Cell Applications) were cultured in an Endothelial Cell Growth Medium (Cell Applications). The culture dish (Farcon) with the cells were incubated in a CO<sub>2</sub> incubator (37°C, 5% CO<sub>2</sub>, in a humid atmosphere). When the cells were grown to be confluent in the dish, 250 µg/ml of bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin was added thereto and the cells were incubated for 33 hours. The mRNA was extracted from the cells by using a FastTack(TM) 2.0 kit (Invitrogen). The labeling of hybridization probe was carried out by using the mRNA according to the same procedure as described above.

[0210] Table 169 shows the expression level of each cDNA in human pulmonary arterial endothelial cells cultured in a medium containing bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin. Genes of which expression was detected in the endothelial cell are as follows:

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45

	HEMBA1000005,	HEMBA1000012,	HEMBA1000020,	HEMBA1000042,	HEMBA1000046,
	HEMBA1000076,	HEMBA1000111.	HEMBA1000150,	HEMBA1000185,	HEMBA1000216,
	HEMBA1000231,	HEMBA1000280,	HEMBA1000282,	HEMBA1000288,	HEMBA1000304,
5	HEMBA1000327,	HEMBA1000338,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366,
	HEMBA1000376,	HEMBA1000387,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000460,	HEMBA1000469,	HEMBA1000501,	HEMBA1000504,	HEMBA1000505.
	HEMBA1000508,	HEMBA1000519,	HEMBA1000534,	HEMBA1000540,	HEMBA1000542,
10	HEMBA1000555,	HEMBA1000557,	HEMBA1000575,	HEMBA1000592,	HEMBA1000604,
10	HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000673,	HEMBA1000682,
	HEMBA1000686,	HEMBA1000702,	HEMBA1000726,	HEMBA1000727,	HEMBA1000749.
	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,	HEMBA1000852.
	HEMBA1000872,	HEMBA1000876,	HEMBA1000918,	HEMBA1000942,	HEMBA1000960,
15	HEMBA1000974,	HEMBA1001020,	HEMBA1001051,	HEMBA1001059,	HEMBA1001060,
	HEMBA1001080,		HEMBA1001088,	HEMBA1001109,	HEMBA1001122,
	HEMBA1001123,	HEMBA1001085, HEMBA1001140,	HEMBA1001033,	HEMBA1001196,	HEMBA1001197.
	HEMBA1001123,	HEMBA1001226,	HEMBA1001112,	HEMBA1001247,	HEMBA1001265,
20	HEMBA1001281,	HEMBA1001286,	HEMBA1001299,	HEMBA1001302,	HEMBA1001326
	HEMBA1001330,	HEMBA1001351,	HEMBA1001233,	HEMBA1001387,	HEMBA1001328,
	HEMBA1001432,	HEMBA1001433,	HEMBA1001435,	HEMBA1001446,	HEMBA1001450,
	HEMBA1001454,	•	•	BA1001510, HEMBA1	•
25	HEMBA1001526,	HEMBA1001533,	HEMBA1001557,	HEMBA1001566,	HEMBA1001569,
	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,	HEMBA1001595,	HEMBA1001608,
	HEMBA1001620.	HEMBA1001640,	HEMBA1001647.	HEMBA1001651,	HEMBA1001655,
	HEMBA1001658,	HEMBA1001678,	HEMBA1001714,	HEMBA1001718,	HEMBA1001723,
	HEMBA1001746,	HEMBA1001784,	HEMBA1001791,	HEMBA1001800,	HEMBA1001804,
30	HEMBA1001809,	HEMBA1001819,	HEMBA1001824,	HEMBA1001844,	HEMBA1001866,
	HEMBA1001888,	HEMBA1001912,	HEMBA1001913,	HEMBA1001918,	HEMBA1001940,
	HEMBA1001960,	HEMBA1001987,	HEMBA1001991,	HEMBA1002003,	HEMBA1002008,
	HEMBA1002049,	HEMBA1002100,	HEMBA1002102,	HEMBA1002113,	HEMBA1002119.
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30	THYR01000085,	THYR01000092,	THYR01000132,	THYR01000163,	THYR01000186,
30	THYRO1000187,	THYRO1000190,	THYR01000197,	THYR01000221,	THYR01000241,
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	THYR01000471,	THYR01000484,	THYR01000569,	THYR01000602,	THYR01000625,
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	THYRO1000988,	THYR01001003,	THYR01001031,	THYR01001093,	THYR01001133,
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40	THYR01001287,	THYR01001320, TI	HYRO1001321, THYF	RO1001374, THYRO1	.001401,
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	THYR01001487,	THYR01001534,	THYR01001537,	THYR01001541,	THYR01001559,
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	Y79AA1000420,	Y79AA1000469,	Y79AA1000480,	Y79AA1000538,	Y79AA1000539,
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Y79AA1002351,	Y79AA1002361, Y7	9AA1002472, Y79A	A1002482	

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[0211] Signal ratios of EC\_AGE\_BSA to EC\_BSA and of EC\_glycated\_BSA to EC\_BSA were calculated for each gene. Genes with high signal ratios were selected. In the case of calculating the ratio of signal value of 40 or less to that of more than 40, such signal values were, for convenience, taken as 40 instead of the real values. When the ratio EC\_AGE\_BSA/EC\_BSA is 2 or more, expression of the genes exhibiting such ratio is expected to be elevated due to advanced glycation endproduct of bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC\_AGE\_BSA/EC\_BSA ranges from 0.5 to 2, expression of the genes exhibiting such ratio is expected to be unaffected due to advanced glycation endproduct of bovine serum albumin. When the ratio EC\_AGE\_BSA/EC\_BSA is less than 0.5, expression of the genes exhibiting such ratio value is expected to be decreased due to advanced glycation endproduct of bovine serum albumin. The lower the value is, the lower the gene expression level is.

[0212] Clone with EC\_AGE\_BSA/EC\_BSA ratio of 2 or higher are as follows: HEMBA1003958, MAMMA1001256, PLACE2000411.

[0213] Clone with EC\_AGE\_BSA/EC\_BSA ratio of 0.5 or less is as follows: MAMMA1001783. These cDNAs are associated with diabetes.

[0214] When the ratio EC\_glycated\_BSA/EC\_BSA is 2 or more, the expression level of the gene exhibiting such ratio is expected to be elevated due to glycated bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC\_glycated\_BSA/EC\_BSA ranges from 0.5 to 2, the expression level of the gene exhibiting such ratio is expected to be unaffected with glycated bovine serum albumin. When the ratio EC\_glycated\_BSA/EC\_BSA is less than 0.5, the expression level of a gene exhibiting such ratio is expected to be decreased due to glycated bovine serum albumin. The lower the value is, the lower the gene expression level is.

25 [0215] Clones with EC\_glycated\_BSA/EC\_BSA ratio of 2 or more are as follows: HEMBA1004850, MAMMA1001256, MAMMA1002132 and PLACE3000119.

[0216] A clone with EC\_glycated\_BSA/EC\_BSA ratio of 0.5 or less is as follows: MAMMA1001783.

[0217] These cDNAs are also associated with diabetes.

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Analysis of genes associated with neural cell differentiation

[0218] Genes involved in neural cell differentiation are useful for treating neurological diseases. It is possible that genes with varying expression levels in response to induction of cellular differentiation in neural cells are associated with neurological diseases.

[0219] A survey was performed for genes of which expression levels are varied in response to induction of differentiation (stimulation by retinoic acid (RA)) in cultured cells of a neural strain, NT2.

[0220] The NT2 cells were treated basically according to supplier's instruction manual. "Undifferentiated NT2 cells" means NT2 cells successively cultured in an Opti-MEM I (GIBCO-BRL; catalog No. 31985) containing 10%(v/v) fetal bovine serum and 1%(v/v) penicillin-streptomycin (GIBCO BRL). "NT2 cells cultured in the presence of retinoic acid" means the cells resulted from transferring undifferentiated NT2 cells into a retinoic acid-containing medium, which consists of D-MEM (GIBCO-BRL; catalog No. 11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin and 10 μM retinoic acid (GIBCO-BRL), and the subsequent successive culture therein for 5 weeks. "NT2 cells that were cultured in the presence of retinoic acid and then further cultured in the presence of cell-division inhibitor added" means NT2 cells resulted from transferring NT2 cells cultured in the presence of retinoic acid for 5 weeks into a cell-division inhibitor-containing medium, which consisted of D-MEM(GIBCO BRL; catalog No.11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin, 10μM retinoic acid, 10 μM FudR (5-fluoro-2' -deoxyuridine: GIBCO BRL), 10 μM Urd (Uridine: GIBCO BRL) and 1 μM araC (Cytosine β-D-Arabinofuranoside: GIBCO BRL), and the subsequence succes-

sive culture for 2 weeks. Each of the cells were treated with trypsin and then harvested. Total RNAs were extracted from the cells by using S.N.A.P.(TM) Total RNA Isolation kit (Invitrogen(r)). The labeling of probe used for hybridization was carried out by using 10 μg of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of stimulation for inducing differentiation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment-of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis, clones with the difference can be statistically detected even when the signals were low. Accordingly, clones with signal value of 40 or less were also assessed for the selection.

[0221] Tables 170-349 show the expression level of each cDNA in undifferentiated NT2 cells, NT2 cells cultured in the presence of RA, and NT2 cells that were cultured in the presence of cell-division inhibitor added.

[0222] Averaged signal values  $(M_1, M_2)$  and sample variances  $(s_1^2, s_2^2)$  were calculated for each gene in each of the cells, and then, the pooled sample variances  $s^2$  were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula:  $t=(M1-M2)/s/(1/3+1/3)^{1/2}$ . When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freeedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information on an increase (+) or decrease (-) in the expression level of a gene in the treated cells when the level is compared with that of untreated undifferentiated cells.

20 [0223] Clones of which expression levels increased by RA are as follows: HEMBA1000005, HEMBA1000042, HEMBA1000046, HEMBA1000076, HEMBA1000111, HEMBA1000141. HEMBA1000282, HEMBA1000304, HEMBA1000307. HEMBA1000338, HEMBA1000150, HEMBA1000185, HEMBA1000387, HEMBA1000392, HEMBA1000428, HEMBA1000456, HEMBA1000376, HEMBA1000357, HEMBA1000508, HEMBA1000519, HEMBA1000540, HEMBA1000469, HEMBA1000504, HEMBA1000459, HEMBA1000545. HEMBA1000563, HEMBA1000568, HEMBA1000575, HEMBA1000588. HEMBA1000557, HEMBA1000592, HEMBA1000604, HEMBA1000622, HEMBA1000655, HEMBA1000673, HEMBA1000682, HEMBA1000769, HEMBA1000774, HEMBA1000791, HEMBA1000726, HEMBA1000727, HEMBA1000749, HEMBA1000943, HEMBA1000822, HEMBA1000872, HEMBA1000876, HEMBA1000910, HEMBA1000942, HEMBA1000991, HEMBA1001008, HEMBA1001020, HEMBA1000960, HEMBA1000972, HEMBA1000974, HEMBA1001085, HEMBA1001043, HEMBA1001051, HEMBA1001060, HEMBA1001071, HEMBA1001077, 30 HEMBA1001122, HEMBA1001140, HEMBA1001172, HEMBA1001094, HEMBA1001109, HEMBA1001121, HEMBA1001226, HEMBA1001235, HEMBA1001265, HEMBA1001281, HEMBA1001294, HEMBA1001299, HEMBA1001319, HEMBA1001323, HEMBA1001330, HEMBA1001351, HEMBA1001361, HEMBA1001377, HEMBA1001398, HEMBA1001432, HEMBA1001435, HEMBA1001442, HEMBA1001388, HEMBA1001391, HEMBA1001454, HEMBA1001455, HEMBA1001517, HEMBA1001497, HEMBA1001569, HEMBA1001570. HEMBA1001718. HEMBA1001723. HEMBA1001581, HEMBA1001585, HEMBA1001620, HEMBA1001711, HEMBA1001864. HEMBA1001869. HEMBA1001761, HEMBA1001815, HEMBA1001819, HEMBA1001861. HEMBA1001967, HEMBA1001964. HEMBA1001888, HEMBA1001915, HEMBA1001918, HEMBA1001940. HEMBA1002008, HEMBA1002022, HEMBA1002039, HEMBA1001987, HEMBA1001991. HEMBA1001979. HEMBA1002084, HEMBA1002102, HEMBA1002113, HEMBA1002144, HEMBA1002160, HEMBA1002049. HEMBA1002226, HEMBA1002229, HEMBA1002267, HEMBA1002162, HEMBA1002185, HEMBA1002212, HEMBA1002270, HEMBA1002337, HEMBA1002381, HEMBA1002458, HEMBA1002477, HEMBA1002508. HEMBA1002590, HEMBA1002645, HEMBA1002628. HEMBA1002558, HEMBA1002561, HEMBA1002583, HEMBA1002850. HEMBA1002712, HEMBA1002728, HEMBA1002780, HEMBA1002661, HEMBA1002678, HEMBA1002935, HEMBA1002951, HEMBA1002968, HEMBA1002939, HEMBA1002886, HEMBA1002934, HEMBA1003021, HEMBA1003033, HEMBA1003034, HEMBA1002970, HEMBA1002973, HEMBA1002999, HEMBA1003064, HEMBA1003067, HEMBA1003078, HEMBA1003086, HEMBA1003096, HEMBA1003129. HEMBA1003142, HEMBA1003148, HEMBA1003166, HEMBA1003175, HEMBA1003197, HEMBA1003199, HEMBA1003202, HEMBA1003204, HEMBA1003212, HEMBA1003235. HEMBA1003250, HEMBA1003273, HEMBA1003276, HEMBA1003278, HEMBA1003291, HEMBA1003309, HEMBA1003322, HEMBA1003328, HEMBA1003348, HEMBA1003369, HEMBA1003376, HEMBA1003384, HEMBA1003395, HEMBA1003463, HEMBA1003597, HEMBA1003480, HEMBA1003531, HEMBA1003548, HEMBA1003591, HEMBA1003595. HEMBA1003640, HEMBA1003645, HEMBA1003617, HEMBA1003621, HEMBA1003622, HEMBA1003637, HEMBA1003646, HEMBA1003656, HEMBA1003679, HEMBA1003692, HEMBA1003715, HEMBA1003720, HEMBA1003725, HEMBA1003729, HEMBA1003758, HEMBA1003803, HEMBA1003805. HEMBA1003836. HEMBA1003838, HEMBA1003879, HEMBA1003885. HEMBA1003893, HEMBA1003908, HEMBA1003937, HEMBA1003942, HEMBA1003953, HEMBA1003958, HEMBA1003959, HEMBA1003978, HEMBA1003987. HEMBA1003989, HEMBA1004000, HEMBA1004011, HEMBA1004012, HEMBA1004015, HEMBA1004024,

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   HEMBA1004049.
                                 HEMBA1004207,
                                               HEMBA1004225,
                                                              HEMBA1004246,
                                                                             HEMBA1004248,
    HEMBA1004199,
                  HEMBA1004200,
                                 HEMBA1004312,
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[0224] Clones of which expression levels decreased by RA are as follows:
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[0225] Clones of which expression levels increase by RA/inhibitor are as follows:
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    [0226] Clones of which expression levels decrease by RA/inhibitor are as follows:
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    [0227] Clones of which expression levels increase in the presence of both RA and RA/inhibitor are as follows:
    HEMBA1000046, HEMBA1000307, HEMBA1000504, HEMBA1000588, HEMBA1000682, HEMBA1000726,
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     Y79AA1001585, Y79AA1001696, Y79AA1001711, Y79AA1002258.
    [0228] Clones of which expression levels decrease in the presence of both RA and RA/inhibitor are as follows:
     HEMBA1000946, HEMBA1003569, HEMBA1005570, NT2RM1000666, NT2RM2000092, NT2RM2000594,
     NT2RM4001754, NT2RM4001905, NT2RP2002047, NT2RP3000980,
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     NT2RP4001950, NT2RP4002408, OVARC1000431, OVARC1001942,
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    PLACE1005923, PLACE1007257, PLACE1010624, Y79AA1000346.
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[0229] These are neurological disease-associated clones.

## Analysis of rheumatoid arthritis-associated genes

[0230] The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)- $\alpha$  participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis.

[0231] A survey was performed for genes of which expression levels are varied in response to TNF- $\alpha$  in the primary cell culture of synovial tissue. The primary cultured cells of the smooth muscle (Cell Applications) were grown to be confluent in a culture dish, and then, human TNF- $\alpha$  (Boehringer-Mannheim) was added at a final concentration of 10 ng/ml thereto. The culture was further continued for 24 hours.

[0232] Total RNA was extracted from the cells by using S.N.A.P.(TM) Total RNA Isolation kit (Invitrogen). The labeling of probe used for hybridization was carried out by using 10 μg of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of TNF stimulation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis,

clones with the difference can be statistically detected even when the signals were low.

Accordingly, clones with signal value of 40 or less were also assessed for the selection.

Table 352 shows the expression level of each cDNA in synovial cells cultured in the absence or presence of TNF. [0233] Averaged signal values  $(M_1, M_2)$  and sample variances  $(s_1^2, s_2^2)$  for each gene were calculated in each of the cells, and then, the pooled sample variances  $s^2$  were obtained from the sample variances of the two types of cells to be compared. The t-values were determined according to the following formula:  $t=(M_1-M_2)/s/(1/3+1/3)^{1/2}$ . When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of

The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the stimulated cells when the level is compared with that of unstimulated cells.

[0234] Clones of which expression levels are elevated by TNF- $\alpha$  are as follows:

55

the gene between the two types of cells at p<0.05 or p<0.01, respectively.

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                                                                  Y79AA1000214,
                                                                                 Y79AA1000346.
    Y79AA1000784, Y79AA1000833, Y79AA1000968, Y79AA1001555, Y79AA1002220.
    [0235] Clones of which expression levels decrease by TNF-\alpha are as follows:
    HEMBA1002150, HEMBB1000240, NT2RM2000469, NT2RM2000984, NT2RM2001688, _ NT2RM4000290,
    NT2RM4000496,
                   NT2RM4000590, NT2RM4001047, NT2RM4001582,
                                                                 NT2RM4001611,
                                                                                NT2RM4001650.
20
    NT2RM4002075,
                   NT2RM4002128,
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                                                  NT2RP1000243,
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    NT2RP1000767.
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                                                  NT2RP2001813,
                                                                 NT2RP2002137,
                                                                                 NT2RP2002928,
    NT2RP2003517, NT2RP2003559,
                                   NT2RP2003564,
                                                  NT2RP2004933,
                                                                 NT2RP2005038,
                                                                                 NT2RP2006365.
    NT2RP3000072, NT2RP3000320,
                                   NT2RP3000484.
                                                  NT2RP3000980,
                                                                 NT2RP3001111,
                                                                                 NT2RP3001420.
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    NT2RP3001495, NT2RP3002056,
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                  NT2RP3003870,
                                   NT2RP3004207,
                                                  NT2RP3004262,
                                                                 NT2RP3004527,
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    NT2RP4000524.
                   NT2RP4000787,
                                   NT2RP4000927,
                                                  NT2RP4000955,
                                                                 NT2RP4000989,
                                                                                 NT2RP4001442,
                   NT2RP4001950,
    NT2RP4001638,
                                  NT2RP4002888.
                                                                 OVARC1001270,
                                                  NT2RP5003524.
                                                                                 PLACE1000246,
30
    PLACE1002816.
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[0236] These are rheumatoid arthritis-associated clones.

Analysis of ultraviolet radiation damage-associated genes

- 35 [0237] It is known that ultraviolet rays give considerably adverse influence on the health. In recent years, there have been significant risks of tissue damage by ultraviolet rays because of destruction of the ozone layer. Thus, ultraviolet radiation has been recognized as a risk factor for skin diseases such as skin cancers (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with skin damage caused by ultraviolet radiation.
  - [0238] After primary cultured skin fibroblast cells were irradiated with ultraviolet ray and were cultured, a survey was performed for genes of which expression levels were varied depending on the irradiation of ultraviolet ray. First, after cultured to be confluent, the primary cultured skin fibroblast cells (Cell Applications) were exposed to 10,000 µJ/cm² of 254-nm ultraviolet light.
- 45 [0239] Messenger RNAs were, then, extracted by using a FastTrack™ 2.0 mRNA Isolation kit (Invitrogen Co.) from the unexposed cells and from the cells that were exposed to the ultraviolet light and then cultured for 4 or 24 hours. The labeling of the hybridization probe was carried out by using a 1.5 μg of each mRNA in the same manner as described above. The data were obtained in triplicate (n=3). The hybridization signals were compared between the cells exposed to the ultraviolet light and the unexposed cells. The comparison was preformed by statistical treatment with two-sample t-test. Clones with significant differences in the signal distribution were selected under the condition of p<0.05. In this analysis, even when the signal is lower than others, the difference in the signal values can be detected statistically. Accordingly, clones with signal value of 40 or lower were also assessed for selection.</p>
  - [0240] Tables 353-509 show the expression of each cDNA in skin-derived fibroblast cells exposed and unexposed to ultraviolet light.
- 55 [0241] Averaged signal values (M<sub>1</sub>, M<sub>2</sub>) and sample variances (s<sub>1</sub><sup>2</sup>, s<sub>2</sub><sup>2</sup>) were calculated for each gene in each of the cells, and then, the pooled sample variances s<sup>2</sup> were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula: t=(M<sub>1</sub>-M<sub>2</sub>)/s/(1/3+1/3)<sup>1/2</sup>. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01

in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the exposed cells in comparison with that of unexposed cells.

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[0242] The expression levels of the following clones were elevated 4 or 24 hours after the ultraviolet irradiation:
    HEMBA1000542, HEMBA1001808, HEMBA1002177, HEMBA1003314, MAMMA1001874, NT2RM2001100,
                                  NT2RP4000657.
                                                OVARC1000004.
                                                                OVARC1001092,
    NT2RP2005732.
                   NT2RP3000592,
                                                                               OVARC1001342.
    PLACE1002816.
                   NT2RM4001002.
                                  NT2RM4001813.
                                                 NT2RM4002266,
                                                                NT2RP2001174.
                                                                               NT2RP2001196,
                                  NT2RP3001216,
                                                 NT2RP3003464,
                                                                               THYRO1000070,
    NT2RP2005358,
                   NT2RP3000690,
                                                                PLACE1006382,
    THYRO1001100, Y79AA1000342.
    [0243] The expression levels of the following clones were decreased 4 or 24 hours after the ultraviolet irradiation:
    HEMBA100005, HEMBA1000150, HEMBA1000156, HEMBA1000158, HEMBA1000168, HEMBA1000231,
    HEMBA1000304, HEMBA1000307,
                                  HEMBA1000333,
                                                 HEMBA1000366,
                                                                HEMBA1000369,
                                                                               HEMBA1000390.
                                                 HEMBA1000464.
    HEMBA1000396. HEMBA1000418.
                                  HEMBA1000434.
                                                                HEMBA1000469.
                                                                               HEMBA1000490.
    HEMBA1000504, HEMBA1000505,
                                  HEMBA1000557,
                                                 HEMBA1000657.
                                                                HEMBA1000673,
                                                                               HEMBA1000682,
    HEMBA1000686,
                  HEMBA1000727,
                                  HEMBA1000752,
                                                 HEMBA1000851,
                                                                HEMBA1000852,
                                                                               HEMBA1000870,
                   HEMBA1001085,
                                  HEMBA1001121,
                                                 HEMBA1001133,
                                                                HEMBA1001235,
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    HEMBA1000872,
    HEMBA1001281, HEMBA1001289,
                                  HEMBA1001299,
                                                 HEMBA1001303,
                                                                HEMBA1001310,
                                                                               HEMBA1001323,
    HEMBA1001595, HEMBA1001620,
                                  HEMBA1001640,
                                                 HEMBA1001678,
                                                                HEMBA1001712,
                                                                               HEMBA1001835,
20
                                                 HEMBA1002321,
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                                                                HEMBA1002341.
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                                                                HEMBA1002999.
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    HEMBA1003684, HEMBA1003799,
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                                                 HEMBA1003989,
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    HEMBA1004133, 'HEMBA1004225,
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                                                                HEMBA1004816, HEMBA1004900,
    HEMBA1005047, HEMBA1005079,
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                                                 HEMBA1005149,
                                                                HEMBA1005152,
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                                  HEMBA1007085,
                                                 HEMBA1007224,
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	[0244] These clo	nes are ultraviolet	radiation damage-as	sociated clones.		

# Table 12

Expression of each cDNA in human tissues (containing clones that are not described in Examples.)

	Clone_name	Heart	Lung	P. gland	Thymus	Brain	Kidney	Liver	Spieen
	GAPOH(Cr1)	38. 210	32.670	23.820	13.580	11.230	21, 120	24.910	22.440
					17.500	92.880	114.650		256.790
	Bactin(Cr2)	279.280		32.749	22.858	26.564	24, 940	22.644	27.001
5	ADRGL 1000005	53. 882	23.005	160. 457	67. 191	101.768	62. 489	67. 150	73.543
3	ADRGL1000007	94.778	85. 185			6.013	5. 219	14. 272	21. 225
	ADRGL 1000009	11.141	50. 520.	10. 357	7. 177			49. 220	22. 221
	ADRGL 1000011	71.656	24. 579	29. 358	19. 473	24.898	30.747		
	ADRGL 1000027	36.238	25. 252	20.855	7. 328	11. 196	14. 298	19.658	11. 288
	ADRGL 1000058	66.209	129.497	55. 226	49.241	30.219	55. 872	67.027	243. 436
	ADRGL1000069	38.630	23.459	28. 991	12.540	27.353	33.633	28. 774	20.911
10	ADRGL 1000077	97.465	63. 656	448. 427	83.412	71.108	53.740	67.906	89. 439
	ADRGL1000092	89. 423	45.692	55, 810	26.033	44. 148	73. 339	96.037	73.091
	ADRGL 1000099	73.675	24.424	36. 128	17.024	25.964	41.391	42.837	29.666
	ADRGL 1000 136	141.745	63.974	77.017	24.777	33.549	58. 986	295.009	84. 985
	ADRGL 1000147	394. 563	155.829	271.210	92.899	165.627	251, 266	253. 420	150. 294
	ADRGL1000159	50.073	25. 425	39. 296	15.194	16.125	20.040	33.720	23. 278
15	ADRGL1000159	69.386	31.051	59. 416	20.154	39.799	27. 027	47.169	20.716
		57.047	23.011	43. 063	23.860	40.581	59.814	117.055	32,630
	ADRGL1000171		18.666	34. 476	15. 434	34.225	32. 962	39, 693	16.334
	ADRGL1000181	45. 892	42. 337	92.865	41.003	45.168	88. 524	85.990	73.392
	BGG111000015	153. 242	94.731	119.688	34.159	30.249	98. 806	98.783	39. 204
	BGG111000016	177.367			20.878	18.769	32, 340	39.666	20.750
20	BGG 11000017	84.712	32.614	38. 131		11.158	18. 241	19, 197	11. 937
	BGG111000022	52.468	20. 452	67.167	12.167	13.313	15, 525	16.757	13, 406
	BGG111000031	30.008	17.072	40.883	12.585	43. 122	43.770	49, 107	38.776
	BGG111000042	49. 926	36.336	51.176	26.964		25, 784	18. 430	25. 385
	BGG111000046	31.618	26. 472	34, 182	31.854	12.650	23, 704	3350. 527.	
	8NGH41000020		2993. 496	1444. 841	537. 162	6973.542	6029. 124 35. 523	37. 596	47.074
25	BNGH41000025	91.717	35. 026	73. 901	27.713	30.765	91.310	75. 797	70.241
	BNGH41000026	176.757	77. 439	98. 345	35. 807	56.991		17.836	23.859
	BNCH41000027	65.029	56. 353	25. 896	22.494	12.763	23.748	72.461	64, 458
	BNGH41000035	148.779	66.776	119. 727	56.576	60.996	96. 959 36. 272	27. 525	24.771
	BNGH41000037	79.500	29.611	43. 438	18.317	20.857	125. 323	86.783	122. 959
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30	BNGH41000048	56.144	32. 253	54.063	14.729	27.312	20. 735	16.100	7.642
	BNCH41000056	67.258	18.694	30.075	15.602	10.072		50.841	45. 285
	BNGH41000087	98. 262	46. 173	77.657	35.329	40.900	50.029	10.725	12.410
	BNGH41000091	50.895	16.985	28. 392	10.147	5.469	22.794	25. 410	24.950
	BNGH41000157	69.043	34, 730	40. 597	18.088	27.072	22.074	25.634	25.843
	BNGH41000169	44.850	21.770	28. 655	11.403	25.991	28.509	15. 553	16. 575
35	BNCH41000181	17. 163	15, 689	13. 948	3.996	9. 287	13, 139	31.695	28. 325
	BNGH41000198	81.510	36.250	60.860	20.585	26.929	35. 751 15. 235	27.285	35. 709
	BNGH41000219	30.302	25. 156	22. 187	13.757	11.208		96.716	78. 266
	BNGH41000229	252.790	65. 948	93. 499	51.108	92.555	101.245 47.456	37.007	39. 131
	BNGH41000237	85.757	46. 997	55. 170	26.780	33.764		39. 238	13. 241
	BNGH41000238	17.744	36.938	42.360	14.922	35.749	42.848 23.918	22.443	27.033
40	BNGH41000243	45. 446	23.667	44. 798	20.875	10.516	12.351	19. 152	22.314
	BNGH41000270	60.889	18.651	29. 618	10.724	15.979	14. 903	32.353	23.777
	BRAWH1000004	43.673	28. 539	7. 640	11.388	19.198	25.732	24.214	24. 767
	BRAWH1000018	59. 409	17. 941	102. 270	17. 107	709.078	55.762	66.754	27. 969
	BRAWH1000021	104.772	29, 951	51.142	21.042	1169.154	70.731	79.670	40. 928
	BRAWH1000027	152, 205	47. 310	67.089	32. 199	64.521	56.487	64.886	100.132
45	BRAWH1000029	106. 376	49. 221	55. 840	40.856	59.552		44 666	1 36 366
	BRAWH1000040	29.419	16.761	31.101	16.622		18. 200	17. 998	69. 194
	BRAWH1000050	161.264	71.786	118. 976	51.863		97.720	81.271 53.856	24. 624
	BRAWH1000051	74.067	34, 341	44.047	20. 726		42.055		
	BRAWH1000060		22. 598	35. 012			38.662	34. 923	11, 200
	BRAWH1000075	17.318	16.898		8.901		17.219		
50	8RAWH1000081	43.025	12.998				17.673		9. 844
	BRAWH1000084		42.178				77.111		102.296
	BRAWH1000095		59, 676				53.814		35.985
	BRAWH1000096						52.880		58.678
	BRAWH1000097		72.506			64. 200			
	BRAWH1000100								
55	BRAWH1000101	134.838	57. 232	106, 632	40.741	96.396	71.642	88. 432	57.336

Table 13

	IRRIWITIANA TATE	Ar 414 T	10 202	14 025	7.695	38.918	23.970	23.794	11.048
	BRAWH1000104	25.414	18.303 5.616	14. 825 12. 463	5.518	6.355	5. 084	9. 107	6.573
	BRAWH1000107	16.949.	492, 704		383.612	368, 156	369,621	277.348	340. 450
5	BRAWH1000110	175.556	68. 459	92.209	45.974	64.703	81.723	90.369	57.301
	BRAWH1000135	199, 303	38. 098	72.093	26.809	57.720	91.568	87,016	35.866
	BRAWH1000190	56.386	41.640	57.914	22.782	55.671	40.034	35. 280	40.134
	HEMBA1000005	11.985	23. 427	18.882	9.766	12.656	9, 959	23. 443	21.677
	HEMBA1000006	37.398	24. 521	24.529	15.587	22.317	13.336	16.038	15. 295
	HEMBA1000012	81.820	57.193	66.828	26.683	55.423	58. 731	85.614	56.259
10	HEMBA1000020	157.967	64.157	115.635	51,940	77.293	77. 321	83. 989	74.362
	HEMBA1000030	82.882	35, 447	66.058	26.464	40.990	60.871	47.058	50.652
	HEMBA1000034	47.434	17.878	50.696	5.594	14.005	6. 673	24.652 37.783	7.134
	HEMBA 1000042	147, 376	94.003	330.908	69.071	76.472	55. 477 22. 788	11.196	15.775
	HEMBA1000045	28. 478	20. 289	20.548	12.445	11.835 68.488	45. 288	37.098	47. 486
15	HEMBA1000046	85.160	84. 475	242.940	57.017	11.185	12. 292	8.491	12.018
15	HEMBA1000047	21.380	18.899	18.166 84.448	11.393	43.131	99. 333	57.041	37. 362
	HEMBA1000048	243. 559	55. 114 11. 876	21.972	7, 477	4.096	13. 675	10, 347	7.770
	HEMBA1000050	22.711 45.071	26.410	38. 158	15. 982	30.754	36. 740	34.184	24.269
	HEMBA1000053 HEMBA1000060	101. 197	34.766	50.643	19.938	34.641	54. 061	42.309	22.530
	HEMBA1000072	240. 166	213.938	224.688	163.030	115.246	207. 809	112.361	276.098
20	HEMBA1000073	23. 202	9. 580	10.815	1.698	6.680	18. 155	12.304	14.973
	HEMBA1000076	95, 997	46. 783	177.931	32.617	48.964	50.792	33.947	44.142
	HEMBA1000084	66.603	25.710	48. 434	18.006	22.553	38.118	40.479	29.683
	HEMBA1000087	70.084	17.515	26.544	8.450	17.590	29. 220	19.519	22.565
	HEMBA1000088	15.474	8.614	19.903	4.775	4.519	11.446	34.905 33.946	6.528 24.614
	HEMBA1000091	80.622	38.604	59.393	23. 956 75. 147	44. 939 54. 384	49.760 70.071	29. 529	55. 422
25	HEMBA1000111	85.814	95. 270	270.542 146.465	37, 419	29. 398	30.694	17.702	30. 398
	HEMBA1000121	55. 476 37. 278	43.368 27.165	34.516	13.619	17, 702	28.069	12.834	23.965
	HEMBA1000128 HEMBA1000129	51.488	19.659	44. 907	12. 208	27.243	30.959	24. 383	26.851
	HEMBA1000141	12.961	24. 515	32. 107	14. 353	13.502	11.152	8. 907	20.635
	HEMBA1000145	29. 273	11,479	20.418	8. 202	9.575	14. 877	10.000	7.817
30	HEMBA1000150	534. 562	326. 814	684.147	211.774	218.448	322. 246	235.752	256.883
	HEMBA1000154	95.272	92.253	101.483	54. 276	42.896	75. 526	92.689	188.019
	HEMBA1000156	50.177	72. 591	58.026	31.149	21.865	38. 964	27.634	50. 220 53. 562
	HEMBA1000158	260.718	63.920	89.680	36. 337	44.915	93. 421 34. 966	44.005	33.564
	HEMBA1000168	74.416	61.152	62.826 28.571	30.512	23.287 19.230	10, 903	11.731	14, 102
	HEMBA1000180	28.502	22. 412 50. 661	213. 994	51.166	43. 435	56. 261	38.862	44. 992
35	HEMBA1000185	21. 302	14.879	16. 948	11. 392	11.821	10.656	12.501	6.979
	HEMBA1000193	14. 122	8.318	11.905	7.519	4.736	3.349	8.544	7.842
	HEMBA1000194	54.688	49. 534	143. 817	37.736	20. 221	34. 328	23.359	56. 497
	HEMBA1000201	21.062	14.098			5. 109	5, 059	9.317	10.522
	HEMBA1000213	22.388	25, 532			17.320	9. 084	8. 469	11.766
40	HEMBA1000216	65.935	51.368			33.659	40. 971	36.328 22.590	34.891 25.781
	HEMBA1000227	52.577	31.332	34.925		18.411	29.778	24. 266	30, 410
	HEMBA1000231	114.369	91, 360				57.000	49, 319	59. 288
	HEMBA1000237	91.024	43. 969			25. 396	32.604	38.910	32, 153
	HEMBA1000243 HEMBA1000244	53.456 173.469	104. 733				124. 532	90. 927	78.610
	HEMBA1000251	22.709	12. 333				13. 221	11.516	11.018
45	HEMBA1000254	74.060					24. 002	20.553	13.215
	HEMBA1000264	29.478	15, 248			3.863	11.228	13.690	3.797
	HEMBA1000269	36, 718	13.465	28.932	20, 412			7.348	24. 793
	HEMBA1000275	66.201	39. 367				49. 267		38.871
	HEMBA1000280	33, 299						21.696 56.916	30.785 92.414
50	HEMBA1000282	93. 815							7.311
50	HEMBA1000287	12. 439							15.510
	HEMBA 1000288	45. 269					_		9. 224
	HEMBA1000290 HEMBA1000296	14.803							17.159
	HENBA1000300	98. 491							
	HENBA1000302	23.840							
5 <b>5</b>									

Table 14

	HEMBA1000303	129. 286	51.013	88.777	32.513	50.462	82.994	44.818	49.271
	HEMBA1000304	112.022	67.470	328.677	54, 678	79. 305	43. 526	38.469	55. 762
	HEMBA1000307	14.054	22.013	31.964	13. 167	15. 571	7.974	10.014	8.685
5	HEMBA1000312	97.082	69.330	183. 923	45. 322	45, 087	52.968	37,741	38.246
			16. 264	18.766	11.688	3.620	10.732	8. 295	14.675
	HENBA 1000318	16.164			41, 731	85.004	48, 526	49, 421	48.866
	HEMBA1000327	29.404	59.618	81. 347				0.392	3.743
	HEMBA 1000333	16.964	13.930	14.530	1.872	5.776	1. 571		
	HEMBA1000338	121.878	62.572	348.751	55. 463	49.114	38. 561	30.698	40.644
_	HEMBA1000343	25. 229	29.781	46.395	20.673	5.872	16.551	10, 139	14.088
10	HEMBA1000349	23,061	12.586	31.755	7.020	17.658	11.622	14.807	15.611
	HEMBA1000351	92.847	57. 338	196.577	41.762	37.094	35. 370	27.645	28.615
	HEMBA1000355	85. 210	38. 388	64, 299	18.101	33.114	43.511	37.808	26.628
		60. 438	38.786	62,442	20.784	17. 894	38.058	40, 431	28.899
	HEMBA1000356			206.803	54. 151	42.793	39. 432	26,076	44.579
	HEMBA1000357	84.898	55.990			20.718	23.499	14.632	23.547
15	HEMBA1000366	47. 131	42.031	90, 450	27.056		34. 985	37.622	36. 900
73	HEMBA1000369	71.428	40.685	54. 384	17.613	21. 422			
	HEMBA1000370	15.354	14.949	22.988	7.916	18. 390	15. 359	13.426	6.647
	HEMBA1000376	80, 183	75. 300	201.705	55.266	66.687	44. 612	55.386	56.070
	HEMBA1000387	100.497	129.367,	351, 196	80.257	104. 250	74.007	57.619	79.876
	HEMBA1000389	69. 342	34.021	71, 118	22.346	27.319	47.936	53.026	34.161
	HEMBA1000390	19. 206	25.788	21.028	12.401	18.372	13.751	16.243	15.036
20		19, 400	22.884	44, 179	8.776	11.742	10. 594	12.266	12.463
	HEMBA1000392 HEMBA1000396	75.409	50. 195	81.870	27.979	30. 393	31, 235	17,771	19.584
					10.341	31.398	31. 214	50,056	18.580
	HEMBA1000411	35.966	24. 397	25. 987			7. 913	6.363	2.306
	HEMBA1000418	8. 165	10.778	14.987	4.031	12.495	44, 729	42.886	34.308
	HEMBA1000422_	93.699	38.329	85.266	39.826	45. 992			
	HEMBA1000428	51.017	30,690	79.229	26.579	24.840	17.767	18. 424	18.608
25	HEMBA1000434	1.747	3.214	11.346	1.210	1.602	2. 927	2.788	2.756
	HEMBA1000442	21.750	7.698	16.227	7.252	3. 336	17.969	11.723	10.645
	HENBA1000443	67, 291	35.910	34.775	26. 420	16.860	31.691	47.856	102.287
	HEMBA1000446	236. 986	69.546	90. 283	32.233	34. 107	119. 377	108.645	60.266
	HEMBA1000456	95. 368	37.560	63, 451	22,640	41.092	65, 256	62.972	43.493
	HEMBA1000459	28. 924	35.333	74.945	20, 475	25. 324	26. 253	13.654	31.317
			27. 246	21.973	9.613	15. 230	14, 091	9.746	16.955
30	HEMBA1000460	18. 649		96. 490	31. 332	83.626	109.503	92.971	62.126
	HEMBA1000452	220. 184	42.636		10.862	15. 595	20, 793	16.716	16.539
	HEMBA 1000464	34. 277	15. 137	27.210				24.048	22.968
	HEMBA1000468	41.755	41.852	68. 356	10.400	23.452	43. 909		
	HEMBA1000469	68.229	71.011	256.705	47. 636	29.853	34. 188	22,568	39.190
	HEMBA1000477	185. 220	47.546	102.939	26.276	40.188	95. 247	52, 454	28.109
35	HEMBA1000481	47.276	37.528	24. 407	17.115	24. 182	29.826	20.717	25.819
00	HEMBA1000488	96.226	31.249	71.522	21.667	27.715	44.499	53.708	33.308
	HEMBA1000490	29.915	13.747	32.568	14.002	12.056	6.900	11.274	7.559
	HEMBA1000491	80.198	22.903	47.786	20.675	32.551	52.682	37.109	28.282
	HEMBA1000498	191.186	112.767	454. 998	88.614	102.997	82.927	53, 205	120.837
		57.318	55. 923	180. 158	44. 170	27.291	34.954	18.532	34, 117
	HEMBA 1000501			7, 152	1.726	0.520	2.245	2,551	1.091
40	HEMBA1000504	1.033	5, 893		21.562	14.691	34.729	19, 508	31.925
	HEMBA1000505	55. 746	36, 631	48, 155		81.505	97. 289	230, 331	95. 150
	HEMBA1000507	204. 165	114, 530	305.249	86.138				
	HEMBA 1000508	205.724	105.067	309, 791	72.709	70.180	77. 388	63.849	45.940
	HEMBA1000518	39. 157	29, 100	31.505	16.650	14.796	15.847	24.729	17.601
	HEMBA1000519	166.937	142.676	468.435	148.478	123.978	128. 646	85.670	111.078
	HEMBA1000520	0.000	0.000	0.000	10.341	10.619	1.488	9.513	9. 395
45	HEMBA1000523	38.708	22.090	40.875	13.852	21.603	32, 384	20.478	21.422
	HEMBA1000531	21.874	34.044	40.027	12.264	11.034	29.775	20.421	12.540
	HEMBA1000534	0.000	0.000			48. 940	25, 365	41.242	72.583
						19.981	17.606	26.698	23.904
	HEMBA1000538	0.000	0.000			27. 059	33.931	16.639	31.893
	HEMBA1000540	21.974	47.343					47,917	
60	HEMBA1000542	64,656	33. 152			35. 278	55.508		47.623
50	HEMBA1000545	148.870	136, 401	48.802		12.534	7, 119	25. 484	15.094
	HEMBA1000547	14.825	20, 199				12.020	13, 535	20. 227
	HEMBA1000551	163.806	171.089				116.646	69, 596	152.516
	HEMBA1000555	10.531	20, 199	25.801	24. 488		15. 431	5, 986	10.933
	HEMBA1000557	80.051	48.396	168.724	37. 150	32.863	31.872	22.800	30. 926
	HEMBA1000561	56. 992	22.797				34.904	24.661	22.470
55	[:	35. 552							
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Table 15 ·

	HEMBA1000563	9. 473	11.545	18.205	6.139	12.689	10.132	7.939	14. 253
	HEMBA1000567	41.385	38. 483	27.881	32. 207	15. 544	26.052	15. 086	86.601
5	HEMBA1000568	44.686	33.379	126.524	26.300	22.533	17. 402	26.970	18.707
	HEMBA1000569	58. 184	27. 187	41,012	21.787	12, 925	36.191	33.944	23. 225
	HEMBA1000575	155, 833	155. 759	434.526	92.140	79, 143	69.949	59.928	71. 189
	HEMBA 1000588	41.087	26.072	31.610	14. 580	18.024	18, 458	23. 553	13. 279
	HEMBA1000590	29.693	17.090	23.618	7.069	6.633	16.725	20.068	13.042
	HEMBA1000591	106, 772	54.874	98.079	34.099	31,776	57, 170	48. 488	32.766
	HEMBA1000592	7, 408	10.031	9. 435	9. 551	8, 209	5. 142	7.480	10.319
10	HEMBA1000594	18, 401	11.048	22.547	15. 327	9. 596	12.099	8.751	6.852
	HEMBA1000594	96, 047	78. 462	146.030	49, 571	36.099	70. 815	41.797	47.748
				46.037				17.317	
	HEMBA1000607	46.819	15.606		9. 438	19, 149	21.038		25. 404
	HEMBA1000608	8, 985	3.040	6.705	0.000	7. 378	4. 453	0.000	5. 544
	HEMBA1000622	45, 570	55.746	113.666	40.310	18. 167	19.390	15, 895	29, 149
15	HEMBA 1000634	126. 532	49.146	138.073	29. 094	95.787	79.662	60, 271	71.657
	HEMBA 1 0 0 0 6 3 6	151.899	51.270	126.200	39.161	51.864	62.611	54.056	39. 415
	HEMBA 1000637	33. 241	23.587	39.380	18.047	16. 265	30. 075	28. 226	24. 559
	HEMBA 1000655	80.165	70.766	219.283	58. 901	61.320	45. 821	40.741	62. 639
	HEMBA 1000557	60,961	31.9931	41.401	18.008	30.565	35. 201	35, 611	42.178
	HEMBA1000662	8,600	8. 490	11.263	5. 475	2, 201	6.140	1.557	2. 504
	HEMBA1000664	14, 358	5.082	3.637	2.670	3.516	4, 913	3.094	3, 579
20	HEMBA1000671	11.588	15. 473	26.067	17. 940	8.865	7.647	10,779	21. 196
	HEMBA1000673	73, 174	77.410	193.253	46. 051	34. 388	33. 975	25.896	31.646
	HEMBA1000675	7.666	12.047	22. 123	5.764	42.036	15. 788	10. 254	15. 555
				21.083					
	HEMBA1000678	7. 453	12.314		12.174	14.897	12.628	6.969	6. 584
	HEMBA1000682	118.965	125.696	255.731	86.894	61.443	66. 299	49.060	82.939
25	HEMBA1000686	25. 079	17. 463	23.126	12.722	10. 282	13.835	21. 393	18. 154
	HEMBA1000702	206.683	94. 357	266.585	62. 386	79.930	90. 914	98. 397	60.559
	HEMBA 1000705	25. 430	25.862	47.190	13.191	19,599	26, 364	25.013	18.833
	HEMBA1000713	56.893	25. 288	70.751	17.660	24. 138	23. 311	21.805	21.736
•	HEMBA 1000718	50.149	43.869	128.515	28. 289	23.213	18.458	10.003	17. 419
	HEMBA1000719	37.969	17.467	28.513	12.147	12.768	22.643	14.744	14. 432
	HEMBA1000722	15, 150	9.762	14.699	6.768	11.726	12.080	5. 907	9. 953
30	HEMBA1000726	159.817	111.276	463.937	91.448	109.093	58. 587	46.517	70.087
	HEMBA1000727	22.867	26.803	28.886	21.475	11, 199	14.966	8.634	30. 401
	HEMBA1000732	28, 630	11.011	12.790	4.617	3. 548	13.325	19.978	13.472
	HEMBA1000736	24. 568	21.982	21.410	7.431	11.376	41,026	31.698	16.801
	HEMBA1000743	0.741	4. 467	1.793	1.637	1. 227	3, 642	4. 563	3.368
	HEMBA1000745	8. 930	7.067	14.546	3.314	10.067	5.403	9. 225	6.085
35	HEMBA1000747	21. 442	12.487	25.662	17.081	5. 384	10. 287	9.865	8. 267
	HEMBA1000748	22. 924	14. 885	35.721	12.634	3.045	11.508	4. 110	11.756
	HEMBA1000749	67. 267	50.826	159.211	43.879	20.345	29.613	19.447	31.693
	HEMBA1000752	54. 929	35.778	162.005	28. 209	31.540	25. 132	15.650	20. 776
	HEMBA1000753	120.889	83.878	155.892	48.092	54. 307	53. 238	38. 941	39. 331
40	HEMBA1000757	20. 234	22. 592	52.608	29.935	23.071	24, 503	14.548	43.779
40	HEMBA1000760	12. 599	38.665	19.973	15.800	30.188	14. 155	10.570	39, 229
	HEMBA1000769	114. 956	74. 924	304.424	86.815	39, 365	48, 405	39.918	55, 931
	HEMBA1000773	2. 162	5. 360	11.883	4. 445	0.965	3.158	3.956	2.663
	HEMBA1000774	128.563	115.732	330.111	84.461	69.618	59. 363	42.656	56, 152
	HEMBA1000780	6.850	7.130	24.176	6.924	Б. 903	6.545	6.667	9. 576
	HEMBX1000783	8. 127	5.076	13.701	3. 276	8.863	6. 241	5. 435	4. 429
45	HEMBA1000791	41.433	51.548	108.542	29.633	42.735	44. 515	43.187	40.856
	HEMBA1000793	108.761	30.885	54.568	18.670	31.512	54.669	45.458	34.788
	HEMBA1000802	15.062	11.125	9.052	10.300	11, 505	12, 950	15.354	16.952
	HEMBA1000813	106.763	52.683	69.701	32, 507	44. 369	65.862	59.842	56.799
	HEMBA1000817	19. 480	7.070	17.915	4.016	15. 239	18. 434	11.273	8.079
	HEMBA1000822	9. 520	10.358	15.760	7.218	8.704	11.185	6.639	4.662
50	HEMBA1000827	96.001	12. 420	24.041	8, 305	24.000	6.709	3. 488	8. 591
	HEMBA1000833	53.675	28. 970	35.897	14,604	26. 383	29.036	20. 591	14. 341
	HENBA1000835		67.353				61, 145		
	HEMBA1000843	74.696		83.737	34.349	42.834	98, 943	66.784	52.015
		74. 227	54. 197	92.042	37.825	58. 571		87.569	55. 077
	HEMBA1000851	23. 913	14.070	13.081	6.847	8.634	12.419	19.200	22. 286
55	HEMBA1000852	56.702	54.074	105.085	31.127	34.200	31.843	28, 843	30. 311
<i>33</i>	HEMBA1000867	15. \$48	10.247	11.912	6.256	1. 227	12.374	8.518	5. 811

Table 16

	HEMBA1000869	19.696	18, 785	34.039	15.061	6. 930	13. 298	14.305	14.712
	HEMBA1000870	64. 189	38. 246	44. 665	12.647	23.970	41.195	21.911	17.508
•				86. 933	36.087	40.608	42.532	43.479	36.141
5	HEMBA1000872	46.848	46.546		11.297	35.077	29. 781	19.453	23. 540
	HEMBA1000875	35. 460	41.166	32.238				36.331	40. 884
	HEMBA1000876	89.976	56.654	194.869	42.595	57.670	53.567		
	HEMBA1000907	22.959	9.656	10.917	3. 599	3. 363	5. 327	13.032	10.676
	HEMBA1000908	45, 409	18.456	30. 665	12.448	8. 174	19.529	24.789	16.299
	HEMBA1000910	47.107	13.681	26. 933	5.866	7.073	19. 938	22.971	11.592
4.0	HEMBA1000918	67. 437	29,880	114.873	25. 206	16.670	25.895	26.769	24.710
10	HEMBA1000919	44. 938	29.704	40. 184	22.126	16.008	24.639	23.073	20. 233
	HEMBA1000934	162.546	35, 314	59.012	18.820	30.796	53. 492	33.824	20.798
		16. 284	29. 481	71.669	12.587	23.834	13.188	7.830	13.322
	HEMBA1000935		39. 296	75.619	25, 080	28.113	39. 401	25.948	30.168
	HEMBA 1000940	44. 243		260.912	62.657	49.118	47. 891	35, 814	49.631
	HENBA1000942	126.095	96.812			8. 391	11.856	11.039	7.414
15	HEMBA1000943	14. 439	12.702	14.690	4. 792	5, 772	0.000	19.405	9. 939
	HEMBA1000946	15.461	5, 506	18.692	9.000				
	HEMBA1000960	179.860	151.073	343.747	107.319	85.691	117.093	82.928	94. 494
	HEMBA1000962	73.395	34.803	60.061	26.562	28. 789	47.944	60.067	31.619
	HEMBA1000968	14. 529	12.486	35. 270	18.733	6. 213	7. 458	7.214	4.624
	HEMBA1000971	50.148	19. 281	37. 515	12. 222	19.562	29.874	22.045	23. 135
00	HEMBA1000972	51.100	33.450	188. 137	28.972	24. 576	23. 736	13.731	27.272
20	HEMBA1000974	5, 609	10.649	12.866	2.929	2, 603	3.800	6.104	4.964
	HENBA1000975	34, 417	19, 132	42, 499	15.644	4.009	15.478	14.192	14.353
	HEMBA1000979	90.061	38. 532	99.641	19:754	27.516	38.801	31.347	36.440
	HEMBA1000981	35. 338	31.281	38.672	19.544	34. 385	38. 280	24.897	29.059
	HEM8A1000983	71.391	34, 501	58.683	22.640	32.825	32. 384	27.465	31.286
	HEMBA1000985	9. 290	20, 363	22.497	4.058	6, 343	9. 035	7.852	3.257
25	HENBA1000986	128.714	74.713	236.019	56. 662	52.957	85. 340	63.718	54.892
	HEMBA1000991	72.707	55.780	160.717	34.676	32.494	41. 317	23.483	37.846
			42.563	69. 807	23.525	34, 263	47.777	47.496	48.154
	HEMBA1001007	123.690		83.746	18.125	25. 490	52. 693	30.668	24.961
	HEMBA1001008	124.864	47.842			17.115	17. 937	17.701	22.055
	HEMBA1001009	37.843	29.269	36.715	11.055		94. 424	47.095	74. 625
30	HEMBA1001014	109.049	83. 356	233. 234	60.123	61.977		27.876	24. 283
30	HEMBA1001017	50, 408	20, 212	48.394	16.020	28.537	31.917		7.188
	HEMBA1001019	7. 327	7. 582	14.865	6.154	10.598	5.643	3.920	
	HEMBA1001020	53.067	55. 646	115.814	31.640	25.647	24. 596	23.146	27. 169
	HEMBA1001021_	115.724	42, 415	59.434	28.828	26.181	64. 484	64.173	29.614
	HEMBA1001022	37.883	25. 835	28.969	18.452	20.270	22.790	25.194	20.783
	HEMBA1001024	23.524	15, 235	16.511	8.023	11,818	13.894	8.606	8.098
35	HEMBA1001026	21.343	12.515	18.851	6.888	7.288	12.663	8.419	7.418
	HEMBA1001043	10.374	11.995	9.892	10.750	19.163	9, 299	8.047	8.589
	HEMBA1001051	124.869	115, 181	387. 345	100.376	67.510	61,660	46. 295	68.994
	HEMBA1001052	38.892	13.860	19.067	12.855	11.445	24. 382	15.726	12.323
	HEMBA1001059	98.097	41.525	66.565	27.826	26.220	46.725	42.356	36.506
	HEMBA1001060	116, 857	74, 020	161.485	61.750	50.524	52, 957	38.575	52.612
40	HEMBA1001064	32. 251	24. 026	33, 937	-14.007	7, 907	13.710	17.387	16.720
	HEMBA1001071	25.850	16.043	19, 924	7.855	3.425	9. 530	6.779	24. 242
	HEMBA1001077	24.689	23, 055	64.486	19.413	16.821	16.858	13.165	12.873
	HEMBA1001078	33.254	26.761	41.713	26, 498	24.531	31, 498	25.302	23.636
	HEMBA1001080	57.701	23. 951	31.254	22.489	24.848	33, 265	31.880	26.484
	HEMBA1001084	62.698	41.625	171.096	31.438	31.760	24, 829	17.487	26.581
45	HEMBA1001085	159. 252	116. 909	294. 247	77.235	81.384	76.498	59.989	55.574
40			42. 537	46.695	19.266	25, 146	33.498	44.927	26.310
	HEMBA1001088	74.704				14.610	11.033	15.558	22.531
	HEMBA1001093	30.048	28.810	72.081	20.831		4, 521	4.834	4.468
	HEMBA1001094	5. 535	8.779	10.059	3.089	4, 628	16,055		10. 190
	HEMBA1001099	18. 322	24. 021	14.814		13.778		11.044	
	HEM8A1001104	21.919	13.788	35.048		18.058	24. 450	21.559	18. 527
50	HEMBA1001109	186.384	190.240	540.908		134.630	93. 324	78.690	116.187
	HEMBA1001114	89. 023	252.529	187.547	75.857	35, 109	66. 259	69.432	341. 702
	HEMBA1001121	32.820	25. 812	89.860	19.710	34. 244	18. 209	9.519	15.621
	HEMBA1001122	3.304	6, 213	8.316	4.763	19, 120	5, 650	4.506	23.059
	HEMBA1001123	108.859	55. 807	190.789	41.415	39.028	42. 683	25.551	30.174
	HEM8A1001133	50.744		36.786	14.764	34.752	25.702	23.524	11.367
55 .	HEMBA1001137	38.685				13.174	15.867	11.767	25.508
	[::C=0/1001131	33.000	1 21.000	1					

Table 17

	LUCUPA 1001146	50 4E2	66, 122	169.353	48.837	60.363	44. 403	30.367	43.561
	HEMBA1001140 HEMBA1001144	60.453			07.291	166.089	101, 134	106.337	142.120
				206.084	46, 342	39.316	61.827	91.170	66.852
5	HEMBA1001145			43.877	13. 337	24.176	19. 965	18.089	28.622
	HEMBA1001158	29.417	28. 121	213.708	37.115	24.460	26. 620	19.178	32.709
	HEMBA1001172	74.727		8.831	7.914	2. 574	8.031	3, 119	4. 980
	HEMBA1001174	6. 279	8.617	43.568	19.819	34. 829	16.588	19.883	17.824
	HEMBA1001175	29. 561	34.909	105. 979	44.066	83. 417	86.736	126.297	79.785
•	HEMBA1001182	136.762	64.608	22.060	9.016	11.018	10. 205	6.347	9.176
10	HEMBA1001184	16.758	9.703	11.626	6.559	5. 738	3. 435	9.089	11.273
	HEMBA1001192	15.119	10.798	110.687	83.431	56. 396	68, 797	99.959	173.379
	HEMBA1001197	82.571	114.743	37. 220	19.514	11.451	24, 172	27.637	12.469
	HEMBA1001208	40.250	30.964 37.345	57.618	18. 958	24.480	52. 160	51.978	31.326
	HEMBA1001213	81.501	16.011	20. 958	17.612	12.418	20. 697	19.108	21.328
	HEMBA1001214	36.798 14.108	10.456	11. 382	7.001	17.058	10. 307	7. 980	11, 111
15		13. 961	14.077	13.384	5. 925	5.876	13.456	12.076	5. 825
	HEMBA1001225 HEMBA1001226	173. 501	137.685	444.754	120.060	113. 306	75, 167	63.960	67.304
	HEMBA1001228	115. 971	48.677	102.518	36.755	64.214	50.002	60.915	35. 178
	HEMBA1001229	246.802	111.161,	135.886	43, 460	94.703	148. 387	156.871	115.302
	HEMBA1001235	43.880	86. 102	81.818	36.759	54. 172	65.830	70.065	66.201
	HEMBA1001238	67.342	62.561	136. 273	36, 471	33.652	41.838	26.195	28.747
20	HEMBA1001242	55. 562	43.106	58.593	41.382	47. 200	38, 498	43.114	44.230
	HEMBA1001247	28.768	22.129	16.518	10.576	8.758	17.031	9.651	13.385
	HEMBA1001253	58.130	60.415	66.640	18.982	45. 992	54.071	95.073	63.393
	HEMBA1001257	33. 557	18.509	24. 256	10.657	12.732	31.261	24.849	9.134
	HEMBA1001261	585.214	143.415	243.791	98.186	169. 988	310.109	234. 388	125.796
	HEMBA1001262	27.336	17. 339	19.088	5. 647	15.678	20.899	11.464	19.889
25	HEMBA1001265	36.604	28.090	152.221	27.730	49.893	34. 423	16.502	26.993
	HEMBA1001266	69.367	67.414	170.657	45.898	31.802	39.554	41.287	52.480
	HEMBA1001269	69.921	44.649	36.964	34. 126	22. 232	42. 207	49.848	39.719
	HEMBA1001272	20.406	15. 416	11.514	7.843	8.604	7.893	20.960	13.545
	HEMBA1001279	113. 597	76.085	147. 371	41.113	50.841	58. 248	43.344	47. 548 56. 625
	HEMBA1001281	45. 326	37.551	65. 225	44.536	46.787	41.371	32.229 196.233	117.566
30	HEMBA1001286	370.697	150. 949	236.623	103.571	123.976	219.461	26.637	19.344
	HEMBA1001289	41.041	24.670	40.151	15.175	30.612	27.627 55.355	46.692	35. 972
	HEMBA1001291	76.537	40.444	50. 226	18.776	38. 423 20. 735	29. 333	17.711	34. 443
	HEMBA1001294	82.258	72.319	157.642	10. 275	15.918	21, 120	15.842	13.595
	HEMBA1001296	53. 487	17.150	31.045	11.631	14, 701	4. 543	9. 800	8. 121
0.5	HEMBA1001297	13.397	24.306 135.140	19.513 326.747	90.817	73.749	56. 152	49.803	80.999
35	HEMBA1001299	122.378 56.839	29.036	56.412	19. 108	20.078	34. 481	51.929	37.087
	HEMBA1001302	14. 975	18. 442	43.778	16.797	10.985	11.442	9.787	19.264
	HEMBA1001303 HEMBA1001306	262.869	135.864	244.234	109.949	109. 582	147. 334	146.509	115.543
	HEMBA1001308	174.017	96.705	220.049	56.953	61.486	74. 225	56.171	58.657
	HEMBA1001310	103.029	52.915	67.714	22.895	38, 245	67. 233	49.204	51.006
40	HEMBA1001312	98.664	47.333	61.080	18.118	33.555	47.007	41.795	38. 527
70	HEMBA1001319	2.396	8.234	13.960	1.828	5. 485	3.003	5.682	3.780
	HEMBA1001322	139.794	39.912	105.709	27.700	41. 977	70.428	70,602	46.470
	HEMBA1001323	33. 347	16.728	25.356	11.399	17. 982	11.181	6.356	12.033
	HEMBA1001326	86.190	37. 984	69.933	24. 331	30.078	49, 223	46.365	16.347
	HEMBA1001327	7. 232	9. 387	23.180	7.314	5. 185	9. 563	4. 423	5. 267
45	HEMBA1001330	115.768	106.951	275.315	73.389	24,661	70.535	40.088	77.680
	HEMBA1001348	15.770	21.874	25.347	9.575	13.666	23. 703	12.647	13.724
	HEMBA1001350	75. 857	38.749	51.454		34. 291	56, 400	34.055	24.753
	HEMBA1001351	52.274	55.313	56.544		46.408	29. 604	52.789	30. 972 29. 131
	HEMBA1001352	68. 321	46.617	54. 427		29. 887	39. 484	29.741	
	HEMBA1001353	39.891	57.492	54.971		27.945			59. 101
50	HEMBA1001358	45.659	52.406	59.774					15. 914
	HEMBA1001361	22. 908	16.519						
	HEMBA1001364	18.896	17.205 22.179	23. 355 38. 795					
	HEMBA1001375 HEMBA1001377	140, 430	131.029						
	HEMBA1001383	23. 974		28.704					
	HEMBA1001387	58. 343							
55	(UCWOVIDO 1301	30.343	1 4. 100			1			

Table 18

	CIENSALAA T	10 000 1	04 C00 I	39, 877	18. 958	10.634	32,922	22.224	33.218
	HEMBA1001388	48.601	24.690		56.412	64. 490	47, 169	44. 169	57.372
	HEMBA1001390	132.003							5. 544
5	HEMBA1001391	18. 302	9.686	12.994	6.299	10.600	8.500	7.116	
•	HEM8A1001398	91.232	50. 992	142, 408	36.081	29. 548	29, 490	28.704	29. 984
	HEMBA1001405	58.645	22. 354	32. 227	15.864	9. 285	19.993	24.564	13.964
	HEMBA1001406	36. 434	22.693	105. 808	18.094	19.994	13.316	18.019	16.592
	HEM8A1001407	38.781	19.637	24, 599	18.935	13. 107	23.014	18.826	15.060
	HEMBA1001411	28. 412	7, 180	21,950	8, 303	9.708	14. 302	8.598	6.663
		66.736	26. 480	35. 635	15.400	24.013	18.356	24.304	20.769
10	HEMBA1001413			18, 414	12. 522	9. 722	12.903	18.283	18. 581
	HEMBA1001414	20.720	7.567			20. 989	32. 235	21.694	26.676
	HEMBA1001415	76.802	54, 702	159.510	34. 156		24. 993	25.847	28, 651
	HEMBA1001416	41.784	23. 474	29. 453	12. 230	24. 881		66.751	
	HEMBA1001432	74.066	60.077	190.870	40.409	63.619	36.879		33.675
	HEMBA1001433	132.672	110. 163	246. 542	77.852	61.676	50. 447	37.821	64. 403
15	HEMBA1001435	138.669	108, 645	334. 104	89. 523	68.855	59.723	58.393	56. 483
13	HEMBA1001442	13.093	8, 604	11, 177	7.985	15.704	7. 291	6.742	6.336
	HEMBA1001446	102.450	63. 255	146.442	40.086	27. 976	37. 353	30.266	41.647
	HEMBA1001450	72. 339	35, 494	55. 103	30.799	31.322	42. 457	42.764	41.349
	HEMBA1001454	146.726	128.060	438. 247	88.679	43.129	54.712	41.131	31.250
	HEMBA1001455	5. 879	8. 197	8, 325	5. 561	4, 437	5, 252	4.300	7.359
		17. 432	15. 927	16.490	6.749	2.733	5, 888	7.836	10.963
20	HEMBA1001459			57. 136	38.874	24.764	19. 473	23.241	32.318
	HEMBA1001461	61.531	52.734	16, 843	12. 984	13.465	48. 381	7.061	25. 992
	HEMBA1001462	10.875	14. 911			51.866	61.784	37.705	68.960
	HEMBA1001463	137. 907	83.753	340.496	93.114		77, 440	27.033	29. 537
	HEMBA1001469	85.416	21, 757	29. 463	15. 911	84. 887		21.371	19.890
	HEMBA1001473	20. 582	31. 855	36.498	8. 307	3.680	16.703		
05	HEMBA1001476	135. 720	113.851	246.800	65. 595	57. 431	63.903	65.229	67.697
25	HEMBA1001477	5. 228	2.001	4. 505	2.645	1.540	3. 243	1.426	2.876
	HEMBA1001478	14. 335	10, 180	12,692	5. 468	4.474	5. 444	2.171	4. 539
	HEMBA1001480	88.891	28. 381	49.689	21.660	14. 126	36, 334	38. 272	30.563
	HEMBA1001483	29.872	5, 156	20. 900	4.647	5, 264	9. 545	13.805	4. 424
	HEMBA1001490	6.867	6. 967	14, 148	7.289	1. 585	5.016	5.792	5. 999
	HENBA1001495	431.282	118.073	203.714	73.985	176.836	195. 947	194.164	146.945
30	HEMBA1001497	93.817	60. 807	227.867	55. 576	41.006	34. 182	23.206	45. 223
		174. 254	120. 414	343. 336	76.008	76. 932	73.234	61.531	76.899
	HEMBA1001510		26. 337	67. 169	15.756	15.962	10.664	9, 567	12.346
	HENBA1001515	45. 158	47, 728	80. 287	34. 595	28. 246	21.020	17.229	33.972
	HEMBA1001517	51.005		7.032	7.566	5. 011	6.466	6.447	4.824
	HEMBA1001522	7.431	8. 980			22, 395	22.767	23.530	17.914
35	HEMBA1001526	48.774	21.300	32.732	18.831		44. 926	26.457	37.421
33	HEMBA1001533	129, 423	85. 570	262.800	70. 163	46.649	53. 508	108.861	25. 371
	HEMBA1001547	59.442	26.656	27.947	8.053	15.558			
	HEMBA1001552	41.663	33, 242	115, 535	26.222	30, 447	18, 258	21.358	25. 853
	HEMBA1001553	58. 388	75.765	66, 228	32, 264	36. 196	54. 513	64.874	41.905
	HEMBA1001557	182.516	80.827	161.852	69.344	80.644	123. 765	111.732	70.946
	HEMBA1001563	39.649	31.429	85. 246	26.057	12.157	15. 987	10.065	17.083
40	HEMBA1001566	37.835	49.964	108. 284	35.793	23. 255	25.180	21.368	39. 375
	HEMBA1001569	75, 584	44, 631	109.624	35. 487	130, 340	63.130	44.960	55.257
	HEMBA1001570	198, 300	125. 319	444, 153	119.332	74. 267	79.979	64.732	90.896
	HEMBA1001579	103, 128	60.654	48.704	22.469	22.629	67.058	24, 391	34. 300
	HEMBA1001581	153.698	126. 225	312.570	131.687	142.104	91.884	67.267	94.418
	HEMBA1001582	3,551	7.087	15. 302	4.019	8, 190	4.888	4.671	5.144
45	HEMBA1001585	27. 271	18. 375	25, 179	14.108	5. 648	14, 993	7.628	12.297
,,	HEMBA1001589	109.877	22.722	49.216	20. 427	22.904	64.665	57.120	21.314
			62.349	46. 938	34. 447	29. 362	34. 516	45.233	35. 562
	HEMBA1001595	71.600			16.004	6.061	21.932	18.414	23, 101
	HEMBA1001604	41.253	27.004	34, 167	21.276	22.867	22.699	14.094	15.366
	HEMBA1001608	35.073	29. 270	41. 525				42.373	58.652
	HEMBA1001615	556. 575	105, 703	103. 519	47.686	27.311	81.914		
50	HEMBA1001620		29, 972	79. 824	31.924	62.056	54. 423	64. 359	36. 203
	HEMBA1001621	70.036	30.704	53. 807	15.048	19.545	42. 391	33. 266	40.516
	HEMBA1001635	39.932	29.397	35.653	16.214	18.765	19.655		14.095
	HEMBA1001636	73, 726	18.596	35. 798	14. 928	12.865		31.819	22.414
	HENBA1001640		45. 105	79. 588	28. 452	22.449		30.009	43.819
	HEMBA1001647	82, 402	39, 456	75. 907	35.084	25. 220		71.158	46.463
55	HEMBA1001651		66.648		51, 802	112.530	208. 201	178.161	95.640
	1.22041001931	1 330, 331	1 33. 545						

Table 19

	HEMBA1001655	60, 368	18. 983	58. 438	20, 404	25.072	27. 162	29. 260	26.673
	HEMBA1001658	6.754	15. 270	17.542	13.420	5.060	4.800	4.973	4. 979
	HEMBA1001661	87.199	20. 304	32.793	13.066	8.394	24.098	22, 916	24.583
5			20, 830	54.460	12.363	48. 457	86. 024	73.847	21.248
	HEMBA1001665	160.583		17.002	34.999	14. 855	17.849	22.906	29.478
	HEMBA1001670	16. 953	38. 651	29.000	10,798	7.763	13.782	17. 314	12. 393
	HENBA1001672	32.013	18.885	34. 336	38.037	14, 401	17.612	30. 520	43, 461
	HEMBA1001673	38. 188		33.810	5. 390	15, 796	13. 173	20.020	12.830
	HEMBA1001675	25. 652	15. 594	85.397	92.681	131.468	50. 365	47. 230	68. 405
10	HENBA1001676	91.000	54.310		93.889	115.305	80. 843	48. 879	83. 933
	HEMBA1001678	218. 382	128.995	335.408		36. 449	33. 368	38. 495	35, 261
	HEMBA1001680	82. 159	51. 521	165.818	33.978	2.581	2,772	2.146	2. 424
	HEMBA1001681	1.654	0.785	0.840	2.142		61.638	30. 820	52.077
	HEMBA1001684	143. 985	84. 151	377.154	72.850	69.097	4. 572	6. 164	7, 330
	HEMBA1001695	16.068	10. 112	14.571	6.860	4, 930		3.870	3.613
15	HEMBA1001702	26.509	13.637	8.186	8.466	4.041	2.043	24. 540	19.603
10	HEMBA1001709	67. 279	26. 552	35. 845	13.982	21.742	28.610		
	HEMBA1001711	20.072	29. 559	39.037	20.902	21. 639	12.713	14.718	33.127
	HEMBA1001712	80. 448	25. 222	51.628	19. 393	12.482	38.014	39. 474	14.831
	HEMBA1001714	360. 368	ب55. 902	142. 225	33.748	51.048	144.094	124.654	59. 543
	HEMBA1001717	78. 599	137. 380	18.549	12.298	5. 575	38. 589	10.120	6.047
20	HEMBA1001718	51.621	52. 280	151.597	31.305	21.166	29.146	14.075	24.411
20	HEMBA1001723	17.072	13.658	8. 525	5.653	8.811	9. 350	11.097	7. 268
	HEMBA1001731	35. 728	22.781	41.531	15. 151	12.421	15. 292	14.020	16. 584
	HEMBA1001734	52. 546	40. 599	99.556	25.099	24.031	28.537	17. 389	32.936
	HEMBA1001736	177. 269	58, 328	110,046	13.820	58. 955	108.630	91.464	62.571
	HENBA1001741	41. 432	12. 649	29.883	14.886	16.207	10.446	11.420	7.286
	HEMBA1001744	5. 531	6.849	12.961	13.191	14. 151	4. 519	8. 367	8.623
25	HEMBA1001745	41.752	17.786	36. 239	12.476	21.118	23.635	15.410	16.514
	HEMBA1001746	27. 437	14.874	24. 099	8.668	21.929	19. 488	11.306	10.070
	HENBA1001761	93. 148	46. 911	179. 597	28.212	33.421	34. 026	19. 164	25. 901
	HEMBA1001762	55.612	45.069	102, 148	38.307	35. 260	33. 316	21.274	45. 248
	HEMBA1001781	13. 298	21.385	26, 693	6.898	17.098	52. 601	11.768	23.068
	HEMBA1001784	89.965	43.765	70.064	26.575	31.708	50. 347	52. 265	31.618
30	HEMBA1001791	182.379	81.719	171.066	44.628	49. 350	82. 856	58, 215	48. 207
	HEMBA1001794	248. 582	163. 789	153.778	73.632	50. 595	152. 279	178.827	132.329
	HEMBA 100 1800	23. 432	21. 165	27.668	11.281	20.728	24. 910	36.900	22.729 10.079
	HEMBA1001803	17. 343	8.333	22.801	6.620	6.043	7. 560	6.613	35. 851
	HEMBA1001804	109.775	44.797	59. 456	29. 337	34. 849	44. 372	36.696	
	HEMBA1001808	78, 129	23.567	18.056	15.858	23. 507	27. 136	41.552	12.332
· 35	HEMBA 1001809	65. 887	31.733	54. 127	33.314	26.179	35. 618 21. 582	15. 074	19.831
	HEMBA1001811	58. 974	24. 196	37. 583	17.314	16.018 29.944	35. 297	25. 257	24.172
	HEMBA 1001815	71. 286	63.775	155, 707	37.153		16.061	22.304	14.646
	HEMBA1001816	38.494	19.017	16. 797	7.139	5. 598 21, 358	15. 313	14. 917	25.144
	HEMBA1001819	18. 590	21. 371	38. 109	20.938	4.470	3.473	2.999	3.099
_	HEMBA1001820	10.884	9. 530	8.017	3.507 62.121	28. 285	42. 988	38. 222	47.532
40	HEMBA1001822	74, 239	95.719	91.314 301.248	95. 135	67.478	89. 045	64.562	61.114
	HEMBA1001824	155. 543	93.583	25. 753	5.777	19.660	19.809	12.020	10.462
	HEMBA1001835	23.616		230, 213	48.968	42.113	39.652	33.559	40.495
	HEMBA1001844	149. 876	52.023	40, 636	20. 235	5. 196	35. 109	20.186	35.814
	HEMBA1001847	52.045	19, 220	250, 547	53.025	28.022	40.644	33.371	35. 250
	HEMBA1001849	101.048	104. 708		15.808	31.525	42.751	44.306	18.213
45	HEMBA1001850	105. 331	27.032	39.813		4, 454	2. 945	0.995	3. 121
	HEMBA1001861	3, 104	4. 469				40.833	22.588	71.713
	HEMBA 1001862	50. 279	145. 708	102.412 50.378			21. 182	21.031	28. 126
	HENBA1001864	24. 313	31.572				25.041	22.874	21.249
	HEM8A1001866	57.711	54, 190 99, 559				40. 562	22.644	40.891
	HEMBA1001869						54. 584	34. 598	42.631
50	HEMBA1001871	75.011	44. 336					8. 554	
	HEMBA 1001876	34. 287	31.955				11.657	14. 965	18.117
	HEMBATOO1878	17. 361	17.619				35, 414	21.946	17.114
	HEMBA1001879	57.004	22. 429						26.320
	HENBA1001884	68.009	84. 640						
	HEMBA1001886	12.711							33.090
55	HEMBA1001888	63. 251	46.960	103.023	41.706	41.134	(3, 111	1 (1.13)	1 33.030

Table 20

	Deur 1001900	42 002	42.848	42.779	30.112	25.432	24.430	22, 605	26.730
	HEMBA1001890	42.902		44, 103	21.972	17, 708	30, 703	19.628	23.571
	HEMBA1001896	66.448	24.720					17. 455	20.813
5	HEMBA1001899	36. 251	25. 553	24. 121	14. 701	12.301	21.838		
3	HEMBA1001904	54. 904	256.020		243.646	55. 587	234. 548	188.571	526.744
	HEMBA1001910	40. 309	10.865	13.738	11.244	8. 226	15. 367	15.894	13. 300
	HEMBA1001911	35. 962	23. 128	26, 357	25, 151	11.860	24. 224	22.870	18.238
	HEMBA1001912	59. 924	66.966	97.679	51.180	45. 903	33. 336	33.019	40.551
	HEMBA1001913	175. 368	39.664	67.432	33. 132	26.376	63.459	70.607	52.824
	HEMBA1001915	14.756	14.666	30. 224	8. 295	7.629	17.718	6.737	8.522
10		5. 018	8. 961	27. 591	7.538	11.032	8. 265	4. 852	4.772
	HEMBA1001918		8. 444	18. 196	11. 252	12.587	7.417	7.668	2.769
	HEMBA1001921	4. 431		4. 664	1. 422	3. 480	2. 935	1. 127	2.898
	HEMBA1001931	3. 948	0.000				37. 223	29.835	13.058
	HEMBA1001939	94. 821	24.679	81.706	24. 209	16. 592	18.649	13.136	19,614
	HEMBA 1001940	54. 512	33. 931	145. 138	26. 273	27.653			
15	HEMBA 1001942	38. 572	16.710	32.402	18.718	14, 782	25. 435	26.410	16. 143
13	HEMBA1001944	210.898	71. 197	96. 883	48. 156	38. 533	82.132	92.097	74.740
	HEMBA1001945	31,531	17.019	14. 533	10. 175	3.037	17.421	12. 222	11.694
	HEMBA1001950	7. 103	7.424	9.611	3. 281	4, 091	7.632	5.310	4.044
	HEMBA1001951	46.024	19, 234	101.026	19. 207	13. 212	23.714	20.006	19, 402
	HEMBA1001958	44, 554	12.806	35. 277	17. 321	13.181	22.652	28.735	20. 948
	HEMBA1001960	20.513	7. 802	16.888	8.822	2.948	8.825	10.834	12. 935
20	HENBA1001962	4. 367	5. 104	4. 205	2.811	3.031	4.870	2.364	2.994
	HEMBA1001964	35, 944	22. 281	62.761	18.757	6.663	17.775	8.068	8.601
		47. 345	29. 504	42.717	13. 526	22.051	33. 555	23.601	37. 521
	HEMBA1001967			16.732	12.797	5. 919	13.447	10. 355	9. 155
	HEMBA1001979	35. 138	6.478		45. 735	24. 898	26. 381	17. 514	28.891
	HEMBA1001987	60.083	52. 275	190.331		50. 862	50.789	40. 252	54.919
05	HEMBA1001991	111. 286	79.833	276. 566	56. 455				24.474
25	HEMBA1002003	66. 389	23, 989	53.710	17.039	17, 174	30. 547	28. 422	27, 345
	HEMBA1002005	86.885	41. 457	150. 127	33.935	15, 339	24. 541	24. 237	
	HEMBA 1002008	32, 101	25. 375	86.511	18. 349	8.912	7.593	18.519	14. 967
	HEMBA1002018	66, 105	22. 380	36.174	16. 334	21.482	27.922	34.098	27.804
	HEMBA1002022	13.985	8.018	13.490	0.000	2.985	5.730	6.036	1.433
	HEMBA1002029	132.547	305.823	115, 974	144.692	70.087	74.071	37.046	204.730
30	HENBA1002030	17.077	10.337	14. 524	5. 906	8. 466	5. 897	6.258	6.824
	HEMBA1002035	48. 658	12.959	10.324	14. 325	7.176	14.446	14.084	13.506
	HEMBA1002037	16. 343	34. 097	27.567	14. 451	12.568	15.087	13.819	12.140
	HEMBA1002038	68. 477	31.733	91, 391	16. 935	8.370	6.020	17.500	19.367
	HEMBA1002039	15. 944	22.707	17.807	13.914	7.910	3.306	4.716	11.003
	HENBA1002042	41.657	27.877	32.654	21, 111	14.815	10.217	24. 300	22.659
35	HEMBA1002043	149. 364	92. 912	208.642	70, 906	53.861	84.089	81. 242	61.829
55	HENBA1002048	137. 253	29. 889	60.279	19.894	21.605	65. 594	55. 483	30. 137
		98. 417	84. 099	271.170	63. 157	87.434	48. 247	39.557	53.676
	HENBA1002049		19. 194	25.821	11.890	16, 358	16.441	25.376	27. 152
	HEMBA1002053	33.636		39.511	37.518	17, 449	25. 297	28.606	39.067
	HEMBA1002055	67. 115	34. 916		14. 136	1.311	8. 564	4. 481	12.538
	HEMBA1002056	13.684	12.039	16. 129		10.973	11. 179	9.701	8. 124
40	HEMBA1002061	11.815	14.960	29.478	10.168	72.416	39. 904	45. 653	53. 581
	HEMBA1002080	59. 350	80.319	81.497		10.773	13.652	6.835	5. 555
	HEMBA1002084	11. 331	7. 502	15. 981	7. 301			43. 117	14,011
	HEMBA1002085	69.868	62.174	111.196	13.760	19.083	101. 175		
	HEMBA1002092	127.409	33.016	60.924	24. 219	32.654	72. 141	50. 433	27.770
	HEMBA1002098	34. 645	16.695	25. 357	15. 741	15.632	18.082	12.882	20, 451
45	HEMBA1002100	118. 301	90.733	129.453	60. 276	41.079	89.713	44.294	67.352
	HEMBA1002101	57.160	69. 427	106.418	34.067	32.565	38. 238	15. 932	74. 139
	HEMBA1002102	104.746	76.058	178.766	45. 801	50.114	53. 399	40.628	54. 459
	HEMBA1002105	35. 380	25.812	31.300	14, 131	14.867	29.842	22.894	23.960
	HEMBA1002107	62.621	45.738	65. 486	28. 199	31, 808	52.057	163.850	77, 437
	HEMBA1002113	745.018	396. 517	1335.986		369.500	391.825	236.013	348.025
50	HEMBA1002119	35. 812	23, 546	72.351	18, 292	19.991	18. 086	26.533	25.611
50	HEMBA1002125	42.105	14. 033	45. 440		20.474	47.217	28.894	33.563
	HEMBA1002121	84. 269	29. 512				40.381	49.691	37, 106
		37.735	19, 103	27. 034			21.694	20.410	24. 305
	HEMBA1002133						9, 451	7. 795	9, 431
	HEMBA1002139	25. 756	10.925				14. 521	10. 225	10. 190
	HEMBA1002141	20.036	14. 349						39. 262
55	HEMBA1002144	86.896	68. 335	193.756	56.749	45.612	36.918	1 43.020	33. 606

Table 21

	HENBA1002147	135.045	48.848	87. 208	42.412	46.318	67. 257	83.313	45. 988
	HEMBA1002150	347, 113	89.434	182.502	48.715	86.270	215. 282	234. 394	85. 507
	HEMBA1002151	50, 410	19.140	11.868	10. 122	7.938	26. 996	19.485	14.196
5	HEMBA1002153	32. 258	25. 478	35, 746	20. 325	25.638	15. 972	26.019	19.827
		118, 226	31.167	44. 382	21.446	21.743	47.426	40, 620	16.858
	HEM8A1002156		114. 853	336. 241	90.651	71.047	63.857	41, 633	55. 419
	HENBA1002160	166.654	68.019	132.156	42.302	37.035	29.438	49. 436	41.818
	HEMBA1002161	72.851	62. 989	307.464	68. 589	51.141	55. 242	37. 823	54. 952
	HENBA1002162	122.516		64. 932	20, 426	7. 151	0.000	30. 979	32.578
10	HENBA1002163	49. 889	43.602		32.652	19, 476	39. 440	43, 639	52.404
	HEMBA1002164	110.221	59.044	71. 408 768. 834	194.638	171.071	159. 271	134. 442	213. 993
	HEMBA1002166	312.586	256. 137			30.091	50. 202	48. 620	13.648
	HEMBA1002167	139.053	18. 430	69. 693	11.057		29. 478	20. 169	23.078
	HEMBA1002173	137.562	47. 918	197.006	36.833	26.164	28. 153	22. 687	14. 081
	HEN8A1002177	100.895	25. 141	41.676	25. 857	17, 903	37. 856	44. 115	27. 390
15	HEMBA 1002178	102.831	19.500	46.717	13. 290	32. 323		59. 110	64.849
-	HEMBA1002179	55.617	56.403	85. 686	45.680	26.918	60.684	22. 480	32.386
	HEMBA 1002185	85.236	71.958	212.844	43.915	27.049	32.172		18. 236
	HEMBA1002188	79.413	28. 280	31.826	23. 275	21.094	33. 295	36.478	30, 532
	HEMBA1002189	56. 349	70.609,	148.011	47.092	32.460	30. 101	34. 751	44. 348
	HEMBA1002191	149.027	80.765	149, 493	49.599	42. 372	60.095	35. 614	13.566
20	HEMBA1002192	15. 125	24.998	24.821	15. 373	16.495	12.778	5, 075	29. 791
20	HEMBA1002195	57.368	28. 635	52.828	16.254	22.600	31.843	32. 995	
	HEMBA1002196 .	14.884	12.040	36.633	16.632	15.443	16. 808	12.691	17. 451
	HEMBA1002199	24.937	13, 539	27, 878	15.728	17. 426	10.639	19.664	8.927
	HEMBA1002204	9, 525	5, 141	14.869	6.784	4.619	10.508	27.818	9.410
	HEMBA1002208	80.832	44, 154	68.317	68.994	37.453	74.064	81.827	112.820
0.5	HEMBA1002212	8.709	6.241	10.946	9.855	2.602	5.864	5. 366	4.214 21.605
25	HEMBA1002215	36.521	28.098	31.165	19. 157	20.170	17.045	19.124	
	HEMBA1002217	50.834	62.759	64.668	59. 460	28.990	37. 379	29. 963	64.813
	HEMBA1002220	27.731	14. 997	21,655	B. 451	6.409	5.663	1.641	6.714
	HEMBA1002226	91.222	113.507	269, 906	85. 183	68. 283	59.461	56. 996	78.924
	HEMBA1002227	55.957	91.527	79.169	45. 309	54.892	28. 856	14. 142	90.668
	HEMBA1002229	170.518	117, 589	418.739	112. 916	121.703	85. 889	63.450 15.399	25. 185
30	HEMBA1002237	47.252	49, 329	124.721	32.838	24. 807	23. 399	45. 873	70. 581
	HEMBA1002239	103.363	107.010	190.830	54.740	72.381	50.451	62.401	55.767
	HEMBA1002241	70.729	45. 281	81.541	43. 824	30.449	54. 328	18. 391	14. 286
	HEMBA1002253	25. 559	27.877	35.744	16.605	13.851	18.938 7.119	4. 456	3.711
	HENBA1002257	6. 344	5. 787	15.404	4. 338	1.225	23. 928	18.619	17. 988
	HENBA1002259	48. 436	19.578	38. 228	12.875	147.403	112.552	83.057	137. 280
35	HEMBA1002262	271.029	219.564	645.284	192, 491	15,078	28.043	29. 609	27.237
	HEMBA1002265	56. 947	30. 786	32.747	58.776	30.097	53.750	24. 099	29.752
	HEMBA1002267	108. 413	102.522	243.566	20.313	15.579	28. 348	19.144	16.695
	HEMBA1002270	51.540	26.396	27.766	11.608	10.900	25. 959	14. 425	10.031
	HENBA1002286	44. 897	17.027	19.776 34.095	19.879	8.778	26, 451	22. 368	13.907
	HEMBA1002290	46.449	29.289	92.158	43.064	48. 204	56.899	80.872	58.027
40	HEMBA1002302	152. 883	48. 105	19. 492	7. 905	4.038	7.098	5. 307	1.737
	HENBA1002304	6.050	132.737	29. 225	24.612	24.050	42. 355	39.076	37, 573
	HEMBA1002307	504.772	93.620	191.534	46. 814	134. 386	238. 599	265. 167	88.087
		2.858	2. 456	9.670	0. 933	4.715	4.369	5.615	4. 579
	HEMBA1002319 HEMBA1002320	10.783	7. 936	12.646	4.775	10.008	4. 330	5, 128	3.630
	HEMBA1002321	10.743	9. 992	10.165			7, 952	4.048	5.700
45	HEMBA1002328	89. 382	28. 578	41.753				34.722	18.301
	HEMBA1002333	63.542	21.208					33. 449	21.452
	HEMBA1002337	93.059	61.863						34. 456
	HEMBA1002339	354. 195	154. 586						192.502
	HEMBA1002341	116.488	29. 538						28. 278
	HEMBA1002341	6.882							4.606
50	HEMBA1002349	6. 318							3.748
	HEMBA1002353	14. 497							16.760
	HEMBA1002356								28.190
	HEMBA1002357	64. 855							
	HEMBA1002360								
	HEMBA1002363								
55	LICHOVIDOC 362	1 11.443	1 31.104	.   55.276	7 44. 111	1			

Table 22

	[icun11002266 ]	13.435	10.346	9. 534	5, 175	9.470	4.446	10.802	9. 325
	HEMBA1002365		4. 107	11. 054	4, 163	3, 224	9.009	7.477	2.921
	HEMBA1002370	29.997	18. 475	11. 325	15. 862	10.204	18. 275	29. 203	18.856
5	HEMBA1002374	91.498		127. 578	52.056	38. 450	99. 590	90. 190	86. 994
-	HEMBA1002376	186.416	75. 425	63.893	37. 221	23.657	110.374	162, 166	50.770
	HEMBA1002377	81.350	41.908	477.021	137. 908	491.500	90.431	81.778	127.767
	HEMBA1002380	189. 521	137.466		125. 938	88.330	90.756	70. 293	106. 965
	HENBA1002381	195.037	101.891				40. 126	24. 243	12.399
	HEMBA1002384	35. 247	22.319	42. 496	14.694	19.780	18.736	15.497	20.728
10	HEMBA1002389	44. 796	8.467	36. 790	11. 793	9.362		22. 294	23.666
10	HEMBA1002396	101.267	69.467	13.025	16. 553	26. 429	25. 964 21. 219	16. 569	22.818
	HEMBA1002402	75.818	24.148	28. 457	8. 848	9. 913		58. 836	38. 486
	HEMBA1002417	132.807	33.708	84, 436	22.910	38.826	58, 589	19.705	13.013
	HEMBA1002419	75.547	31.202	41.690	13, 558	16.457	27. 281		
	HEMBA1002420	20.818	20.448	35. 559	17.034	13.878	23.652	14. 721	24.637
4.5	HEMBA1002421	23.903	25. 285	59.023	7.957	14. 189	24. 230	61.011	
15	HEMBA1002423	12.762	11.755	25. 941	12. 938	14.177	14. 263	12.495	7. 512
	HEMBA1002424	111.995	32.293	45. 657	24. 424	25.667	42.797	41.513	31. 249
	HEMBA1002426	60.617	23.489	45. 906	20, 305	25. 173	30, 860	37. 738	21.223
	HEMBA100243D	24. 143	3. 128	4. 900	1.517	4.594	3, 316	8. 552	3.069
	HEMBA1002439	59.808	37.475	93.025	15,789	23.324	47.857	33.099	27. 888
	HEMBA1002441	77.869	99. 262	110.341	38.723	34.562	65.309	85. 421	66. 581
20	HEMBA1002454	58. 292	15. 281	38.384	7. 520	19.044	25. 972	22.845	22.015
	HEMBA1002458	57. 329	46, 103	101. 242	30. 906	82. 184	61.800	26.094	59. 039
	HEMBA1002460	32.814	9. 205	25.085	12.160	23.009	18.683	14.678	14. 249
	HEMBA1002462	98.420	38. 135_	55. 208	10.919	24. 257	49.697	43.851	32. 387
	HEMBA1002465	11.819	15.260	28.272	11.939	11.225	10.938	13. 593	20. 635
	HEMBA1002469	129.538	61.348	120. 187	39. 999	39.213	76.320	69.012	86. 309
25	HEMBA1002475	3.180	5, 116	4. 323	2.230	1.467	4. 495	10.058	15. 691
	HEMBA1002477	93.696	64.730	238.114	55. 207	43.349	42.487	29.532	52. 786
	HEMBA1002480	210.023	58.823	84.566	37.478	45.060	106.554	97.791	70. 487
	HEMBA1002481	104. 499	76.474	222. 903	71.502	68.097	67.421	42. 334	82. 875
	HEMBA1002486	81.465	42.269	169. 291	49. 953	40.852	39.475	29. 153	26. 233
	HEMBA1002490	66. 595	11.331	31.314	14. 602	25.852	35. 945	35.954	15. 278
30	HEMBA1002495	59.387	12.315	25. 235	7.937	4.091	17. 402	14. 269	10.773
	HEMBA1002498	56.425	23.969	67.108	11.632	15.655	24.420	8. 272	12.219
	HEMBA1002501	40.955	16.994	22.074	13.575	16.498	21.707	19. 506	24.619
	HEMBA1002503	81.763	65.044	154. 595	39. 638	33.778	31. 214	32.219	26.800
	HEMBA1002504	155. 357	95.219	279.391	90.092	120. 246	70.516	52.190	53. 323
	HEMBA1002508	99.443	88.234	259. 961	107.085	79.039	59, 181	59. 924	61. 423
35	HEMBA1002513	50. 560	22.902	30. 431	26. 184	20.783	30.500	32.903	22.864
	HEMBA1002515	60. 938	23.064	25.098	16.172	5.716	20.264	20.643	13. 727
	HEMBA1002524	94. 350	35.789	56.675	25. 998	28.978	49.840	57.148	25. 205
	HEMBA1002538	116.609	19.632	26.764	12.798	20, 203	16.422	17. 588	15. 759
	HEMBA1002542	81.641	81.952	188.888	54.986	41.864	32.890	30,719	38. 321
	HEMBA 1002544	52.394	49.175	98. 415	47.569	28.375	28.766	20.948	46.753
40	HEM8A1002546	76. 538	62.763	156.051	47.625	74.374	45. 975	34.756	7. 127
	HEMBA1002547	11.448	4. 516	10.647	4. 733	12.220	11.801	9, 959 94, 586	
	HEMBA1002550	67. 373	39. 322	48. 468	15.671	16.497	121,814		25. 401 18. 665
	HEMBA1002551	94. 391	14. 109	27. 085	11.976	8.787	41.811	16.656 63.216	
	HEM8A1002552	204, 583	77. 430	205. 444	49.448	44.756	67.408		57.684 7.416
	HEMBA1002555	25. 583	16.987	6.743	7. 020		14.795	10.111	
45	HEMBA1002558	92.744	77.405	245. 703	59.079		33.253	41.617	27, 124
43	HEMBA1002561	53, 810	51.725	155.895			17. 264	9,056	
	HEMBA1002562	15. 261	10.822				12.036		
	HEMBA1002568	24. 946	17.442				15. 262	16.158	22. 328
	HEMBA 1002569	112.340	34. 133				54. 936	26.164	27. 309
	HEMBA1002570	43. 528	50.809				28.874	9.812	50. 494
50	HEMBA1002574	106.101	25, 148						31.310
50	HEMBA1002583	36.042							16.418
	HEMBA 1002587	61.527						40.250	39. 915
	HEMBA1002590	151.583					58. 221		53. 855
	HEMBA1002592	97.854	85. 949						57.578
	HEMBA 1002595	146.016	25.688						26.760
	HEMBA 1002609	97.442	41.926	56.054	29.427	35,650	35.839	44.688	47.074
55									

Table 23

			** ***	70 446 T	72 277 1	12.909	21.055	16.612	49.136
	HEMBA1002617	26. 792	86.617	59. 446	73. 277			35.003	28. 108
	HEMBA1002619	101. 131	25. 998	30.959	15. 921	21.913	40.814		12. 283
	HEMBA1002621	14, 592	25.845	18.082	8.927	7. 391	8.869	5. 823	
5	HEMBA1002524	254.635	42.837	73.568	48.036	71.673	113. 228	101.786	53. 514
	HEMBA1002628	13.044	21.509	23. 549	9.956	16. 559	10. 257	7.527	11.624
	HEMBA1002629	32.199	16.370	29. 306	15.884	5.722	15.410	42.964	19. 580
		55. 206	48, 044	90, 986	36.904	27.840	28.811	37.912	40.048
	HEMBA1002632			220. 184	68.171	48.643	56.847	41.355	59.667
	HEMBA1002645	95.909			16.958	11.617	23. 904	29.214	16.599
	HEMBA1002651	39.882	27,730	33, 313			37. 216	25.827	23.282
10	HEMBA1002652	107.869	24. 187	46.646	22.248	22. 950		45, 193	
	HEMBA1002659	133. 320	62.916	259.854	57.860	53. 172	46.511		47.291
	HEMBA1002661	88.495	68.014	154, 170	35.196	22. 499	26.290	22.314	23.727
	HEMBA1002666	34, 174	20.511	39. 391	17.036	15.852	20. 842	19. 202	13.470
	HENBA1002667	155, 384	166.244	164.658	29.523	520.013	30. 234	25.612	83.769
	HEMBA1002673	71.650	40.718	73.822	33.403	39. 914	40. 129	38. 519	22.532
15	HEMBA1002678	161.681	89.986	247.534	84.722	54. 176	46. 941	61.944	77.085
, •		56.416	61.838	66. 537	37.679	18.172	29, 420	38.238	44.113
	HEMBA1002679		3. 364	5. 387	3.816	1.793	4. 608	3, 600	2.944
	HEMBA1002688	6.756			14.788	12.463	31.752	34, 100	14.772
	HEMBA1002696	49.639	17.555,	29. 241		50. 688	113. 980	87.727	59.878
	HEMBA1002703	185.328	96.718	97.793	54. 473		26.777	29. 277	29. 224
	HEMBA1002706	49, 533	30.340	35. 679	18.469	19.118			28.457
20	HEMBA1002712	52.878	59.111	110.506	41.591	43. 597	39.604	30, 872	
	HEMBA1002715	149.045	59.858	87.643	47.473	41.264	95. 279	127, 808	65.580
	HEMBA1002716	23. 142	6.155	17.077	15.783	23. 557	19.064	27.647	7.572
	HEMBA1002718	26, 328	19.063	41.749	26.345	16.735	28. 367	26.822	21.779
	HEMBA1002728	117.984	88.950	293.019	81.290	43.679	65. 830	46. 321	57.003
	HEMBA1002730	131,726	26.862	67.877	28.628	36, 686	49. 987	50, 380	43.208
25	HEMBA1002734	77.679	26. 481	34.604	21.128	21.756	41, 413	80.057	45.992
25		10.730	11.276	12.768	7.910	1, 394	8. 502	8.297	10.909
	HEMBA1002742	60, 876	22.803	35. 400	15.830	15,630	30, 605	31.889	32,759
	HEMBA1002746			38.669	17.760	32, 833	43, 493	53.440	49.691
	HEMBA1002748	75.748	26.130			10.037	22.527	29. 331	30.774
	HEMBA 1002750	40.663	45.306	95. 205	18.200		44. 593	28. 497	39.737
	HEMBA1002755	94.758	62.505	220.964	63.414	37. 572	8.714	10. 261	5. 285
30	HEMBA 1002759	13.935	3.117	8.450	3.792	2. 291			
	HEMBA 1002763	430.941	88.931	172.920	71.623	88. 921	195. 471	197.995	118.224
	HEMBA1002767	65. 682	25.272	35.782	14.035	19. 183	31.497	33. 393	18. 347
	HEMBA1002768	100.803	57.554	59. 457	35.570	28.006	43.770	40.930	38.215
	HEMBA 1002769	103.210	30. 236	54.098	17.099	19.753	35.636	41.922	26.940
	HEMBA 1002770	20.350	16.268	28.054	21.736	10.754	12.030	14.991	11.776
	HEMBA 1002777	130.615	37.655	72.072	41.794	31.219	54. 881	59.342	43.652
35	HEMBA1002779	97. 457	29. 259	75.705	22.719	22.643	33.689	38.357	27.804
		72.338	50.411	181.356	42.070	19.957	31, 370	27.642	39.672
	HEMBA1002780	87.371	61.291	152.514	38.033	29.616	28.032	20. 352	34.761
	HEMBA1002790		77.515	95. 182	31.252	41.834	100.167	80, 301	50.036
	HEMBA1002794	202.405			20.281	12.823	11. 156	11.647	15.735
	HEMBA1002798	9.194	21.334	22.468		4. 420	3,016	13.829	6.693
40	HEMBA1002801	10.311	4.603	11.704	3. 190		60. 935	44. 046	51.794
	HEMBA1002810	42.583	45.313	55.088	35.416	29.480	33.389	50, 974	49.045
	HEMBA 1002816	52.084	37.823	56.994	35.902		152.339	171. 186	117, 409
	HEMBA 1002818	321.516	100.826	187.799	84.893	81.695			
	HEMBA 1002820	139. 924	107.278	533. 137	90.533	79.745	59, 869	54. 302	52.958
	HEMBA1002826	40.776	6.495	16.825	5.349	3.319	11.765	7.355	8.363
	HEMBA1002833	119.102	44.248	40.839	17.864	23.748	44. 398	57.302	36.668
45	HEMBA 1002850	5. 941	8.407	13. 251	6.179	2.932	4. 352	4.844	3.735
	HEMBA1002862	60.735	32.524	30.030	9.693	9. 527	27.595	19.397	18.101
	HEMBA1002863	77.126	30, 401	44.872			50.264	55.374	45.005
		25. 385	13. 583	42.122	15. 283		22. 992	15, 180	16. 196
	HEMBA1002867			38.073			53, 318	51.363	
	HEMBA1002876	101.249	55, 603					6. 432	
F0	HEMBA 1002886	9.474	14.188				36.684		
50	HEMBA1002896	78.580	27.420						
	HEMBA1002913	126.001	32.845					56.608	
	HEMBA1002921	63.378	25.443					37.595	
	HEMBA1002924	65.007	29.109						
	HEMBA1002934	432.841	308. 291	644. 522	180.470				
	HEM8A1002935	92.005	52. 184		49, 477	41.867	34.331	29.646	38.889
55									

Table 24

	DICHOL LANGAGE	20 000 1	20 044 1	33.817	12.784	18. 251	14, 107	24. 131	18.662
	HEMBA1002937	38.698	30. 844	33, 838	19.077	13.734	19. 266	17.364	17.750
	HEMBA1002939	39.755	22. 867		21.860	18. 241	23. 920	21.112	16.286
5	HEMBA1002944	53, 762	33. 349	51.861		29.614	19.702	28. 422	21.177
	HEMBA1002951	38.716	29. 783	39.196	19.808	13.758	15.056	7. 297	13.424
	HENBA1002954	24. 907	8. 542	20.941	9. 265	37.753	44.037	31.812	41.725
	HEMBA1002962	86, 680	62.578	220. 246	62.027	40. 380	43.093	38. 816	50.281
	HEMBA1002968	105.871	78.850	221. 414	65. 545		17. 125	23. 514	36. 180
	HEMBA1002970	48.034	34. 741	30.834	18.482	6. 639	25. 193	14. 823	23. 202
10	HEMBA1002971	39, 492	44. 145	35.618	25.614	12. 932	29. 947	26. 101	34.769
••	HEMBA1002973	83.710	70.965	156.167	43.307	28. 902	11. 324	16.059	13. 956
	HEMBA1002978	35.833	19. 362	27. 056	13.075	20. 398	33.054	37, 199	21, 249
	HEMBA1002981	107.112	35, 200	56.576	23.695	26. 105		28. 927	20. 335
	HEMBA1002985	79.217	44. 154	116.532	27.950	26. 158	37. 462	28. 919	20. 529
	HEMBA1002986	61.056	78. 203	68. 834	49.967	64. 529	38. 333	15. 956	22, 796
15	HEMBA 1002 988	37, 307	36,609	71.802	20.621	8. 965	16. 229	61.291	91, 211
13	HEMBA1002992	97.120	72.656	79.841	50.454	34. 289	57.004	19.510	48. 529
	HEMBA1002995	51, 473	63.779	55.081	36.903	25.007	38.630		25. 973
	HEMBA1002997	41.734	70.805	29.264	27.019	33.664	24. 201	18.442	9.970
	HENBA1002999	35.341	16. 456	18. 357	11.146	7, 034	12.086	13.966	20.055
	HEMBA1003004	55, 654	33.689	35. 194	15.119	16. 204	20. 866	27.891	20. 924
20	HEMBA1003006	40.682	24.885	20.750	20.903	26.595	25. 445	20.310 5.688	12.638
20	HEMBA 1003008	29.269	20. 922	74.697	25.061	17.787	10. 271	60.969	90. 296
	HEMBA1003021	130.889	123.646	311.225	101.957	95. 443 52. 131	64. 844 26. 286	28, 112	31.561
	HEMBA1003027	54.935	32.610	44,710	18.890	34, 111	29.704	49. 230	45. 833
	HEMBA1003029	33. 333	42.436	60.787	20.829	13, 316	13.955	15.773	27. 136
	HEMBA1003031	34.000	25. 311	18. 494	14. 998	50. 526	81.278	84.036	46. 352
25	HEMBA1003032	171.114	46.990	71.365	23.640	90. 670	70.150	55.336	77.819
25	HEMBA1003033	168, 563	118.674	378.771	109. 222	85.630	61, 733	36.799	63. 312
	HEMBA1003034	173, 162	127.221	484.135 9.305	108. 238	5, 058	11.024	2.553	4.409
	HEMBA1003035	11.693	5. 195	145.321	58. 521	65. 732	104.677	89.571	71. 574
	HEMBA1003037	261.159	89.481 105.085	291.931	93. 188	75. 193	53. 097	39.564	58. 217
	HEMBA1003041	103. 945	39.965	46, 856	26. 192	11.615	35.659	25.378	32.416
30	HEMBA1003046	127, 888	49.341	139,750	32, 219	32.320	57, 450	33.390	28.702
30	HEMBA1003047		35. 962	42.305	12.040	20.442	39.108	29.597	21, 461
	HEMBA1003048	87. 433 6. 366	8. 535	6.201	8.809	4.415	7. 239	3.330	7.829
	HEMBA1003064 HEMBA1003067	55.833	34. 508	77.097	26. 154	20. 523	28.755	24.783	17.488
	HEMBA1003071	54.728	22.509	28.869	17. 461	19.647	20.624	22.285	19.438
	HENBA1003072	62. 421	30.769	31.225	26. 145	22.906	21.483	17.616	19. 134
35	HEMBA1003076	111. 254	51.085	78.972	37. 151	40.422	49.911	47.023	64.737
33	HEMBA1003077	36. 471	15.407	24.522	8.009	8.453	18.661	13.797	5.837
	HEMBA1003078	34, 143	38.741	77.906	31.907	37.169	17.933	17.439	18. 923
	HEMBA1003079	28. 559	39.563	41.646	26.110	25.889	25. 576	18.026	24. 526
	HEMBA1003083	61.036	48.635	169.439	52.788	60.016	41.611	29.619	67.469
	HEMBA1003086	49.032	40.488	154.409	29.869	12.063	16.544	16.039	19. 219
40	HEMBA1003090	34.778	14.860	23.758	12.710	24. 132	15.848	25.027	14. 265
40	HEMBA1003094	184.999	43.363	72.116	30.096	53.636	78.251	84.551	34.775
	HEMBA 1003096	31.440	18.030	25.774	10.290	11.781	14.033	27. 791	11.348
	HEMBA1003098	36.774	64.970	88.562	34.074	24.271	25.656	18,003	31.059
	HEMBA1003101	55.716	24.121	22.316	11.682	13. 163	21.315	25. 117	15.689
	HEMBA1003109	48, 411	21.093	19.285	21.315	21.724	27.826	31.034	21.809
45	HEMBA1003114	41, 101	24.785	22.792	14.164	14.657	18. 320	15.152	16.038
45	HEMBA1003117	22.939	13.535	20, 191	6.812	10. 538	14.917	18.015	12.566
	HEMBA1003120	24. 531	24.408	55.805	26.574	13.838		15. 080 26. 267	21.728 38.207
	HEMBA1003129	40. 276	46.792		37.995	37. 989	21.990		
	HEMBA1003133	50.080	22.873			20.000	21.592	25. 551	27. 656 25. 346
	HEMBA1003136	146.630	23.706			31.049	69.754		
50	HEMBA1003142	69.008	47.867			30.384	25. 274		29. 493
~~	HEMBA1003148	59. 282	20.084			18. 973	32.206		24. 674
	HENBA1003151	53.856	20.003				27.114		13.546
	HEMBA1003152	20.577	9, 803				31.586		6.853
	HEMBA1003157	16.477	9. 272						10. 181
	HEMBA1003166	293.814						148. 208 5. 733	7. 164
55	HEM8A1003171	17. 730	8.702	16. 527	5.499	6.963	1.301	1.133	1 1.104

Table 25

FEBBA10031718   51.3815   31.8815   30.182   31.7815   32.530   31.0702   37.7715   1.7816   32.530   31.0702   37.7715   1.7816   32.530   31.0702   37.7715   31.8816   32.530   31.0702   37.7715   31.8816   32.530   31.0702   37.7715   31.8816   32.530   31.0702   37.7715   31.8816   32.530   31.0702   37.7715   31.8816   32.530   31.0702   31.8816   32.530   31.5816   32.530					100 100 1	20 004 1	17 624	21 152	13.386	15, 936
TRIBUTION   15   100   45   17   17   17   17   17   17   17   1		HEMBA1003175	38.620	40.445	100. 302	29. 594	17.624	21.152		
FERNALOGISTES   100, 461   75, 611   231,782   75,781   68,278   54,272   55,862   61,615     FERNALOGISTES   15,472   27,537   45,331   20,623   18,400   21,785   30,501   35,864     FERNALOGISTES   15,462   9,564   5,514   5,965   4,051   3,118   7,054   7,064   7,066     FERNALOGISTES   14,695   18,409   81,183   15,696   4,051   3,118   7,054   7,064   7,066     FERNALOGISTES   15,772   17,575   18,409   81,183   15,696   41,200   31,106   32,336   45,183     FERNALOGISTES   15,772   17,575   18,405   13,183   15,696   41,200   31,106   32,336   45,183     FERNALOGISTES   15,773   15,787   47,406   41,200   31,106   32,336   45,183     FERNALOGISTES   15,415   13,105   15,070   5,771   4,792   31,681   10,789   6,536     FERNALOGISTES   15,415   13,105   13,670   5,771   4,792   31,681   10,789   6,536     FERNALOGISTES   15,415   13,105   13,670   6,371   4,792   31,681   10,789   6,536     FERNALOGISTES   15,775   12,184   40,484   11,586   40,780   40,413   9,856   14,985     FERNALOGISTES   15,775   7,015   42,484   11,586   40,770   70,775   55,675   44,677   70,154   13,745     FERNALOGISTES   15,775   7,015   42,493   31,776   70,775   55,675   40,191   73,748     FERNALOGISTES   15,775   7,760   12,586   14,393   17,475   14,707   70,154   13,745     FERNALOGISTES   15,775   7,760   12,586   12,393   47,500   4,815   7,247   70,775   55,675   4,391   4,756   4,467   4,476   4,4		HEMBA1003179	63.835	33.869	50.631	27. 163	25. 502	35.500	39.052	37.713
TREBUTION   18   36-22   27-357   45-533   20-623   8.740   21-755   30-501   35-864   70-66						75 781	58 278	54, 222	55, 862	61.615
REBRITOSTIPS   8, 452   7, 564   7, 534   7, 535   7, 595   4, 051   3, 138   7, 054   7, 066   REBRITOSTIPS   8, 452   7, 138   18, 407   81, 153   15, 156   157   157   99   942   7, 138   15, 157   158   158   1	5									
International Content   1.5		HEMBA1003196_	36.422	27.557						
		HENRA1003197	8. 462	9.564	5. 534	5.955	4.051			
					81 183	15.696	16,799	9, 492	17, 381	15. 917
HEBBAT003279   12,713   35,727   172,818   48,560   31,451   31,191   27,471   28,849   HEBBAT003271   12,334   30,705   372,474   41,164   52,397   59,653   45,714   54,383   HEBBAT003272   11,345   31,050   372,474   74,164   52,397   59,653   45,714   54,383   17,475   18,581   10,186   53,586   53,587										45 183
THE BRANDOSTR   123-113   123-128   135-129   136-15   15-351   19-417   17-503		[HEMBA1003202								
		HEMBA1003204	66.523	56.272	172.818					
				52 768	35, 498	5. 529	38. 451	16.353	59.417	17. 563
	10					74 164	62 392	59.663	45.714	54, 363
HERBAT003220										
FERBATIOD3227   25.003   22.831   28.577   7.996   11.404   10.413   9.856   14.985     FERBATIOD3227   105.736   71.288   40.848   1.585   20.280   48.241   44.574   19.547     FERBATIOD3227   30.394   26.363   41.333   22.958   77.475   14.707   20.154   19.749     FERBATIOD3227   40.394   26.363   41.333   22.958   77.475   14.707   20.154   19.749     FERBATIOD3226   69.643   70.015   42.433   31.176   20.775   55.815   40.191   75.288     FERBATIOD3226   8.677   77.886   8.735   7.270   7.328   77.285   5.285     FERBATIOD3250   7.260   12.5984   10.9526   33.333   19.405   70.834   22.018   23.855     FERBATIOD3250   55.274   51.495   65.197   28.241   33.512   44.917   62.506   60.075     FERBATIOD3255   55.274   51.495   65.197   28.241   33.512   44.917   62.506   60.075     FERBATIOD3275   71.79   16.603   40.414   13.99   19.441   33.983   28.814   19.028     FERBATIOD3276   19.492   18.996   46.948   14.157   12.769   11.524   8.622   77.414     FERBATIOD3277   43.38   73.37   27.27   77.79   77.28   77.28   27.993   77.240   27.704     FERBATIOD3277   35.779   34.802   113.584   23.812   17.208   20.437   14.885   22.485     FERBATIOD3277   35.799   34.802   113.584   23.812   17.208   20.437   14.885   22.485     FERBATIOD3278   35.798   24.905   71.222   17.479   15.791   15.767   10.948   17.401     FERBATIOD3278   35.798   27.905   71.222   17.479   15.791   15.767   10.948   17.401     FERBATIOD3278   35.998   24.905   71.222   17.479   15.791   15.767   10.948   17.401     FERBATIOD3278   35.998   24.905   71.222   17.479   15.791   15.767   10.948   17.401     FERBATIOD3279   13.248   39.383   37.987   37.785   37.991   39.131   39.131   39.383		HEMBA1003218	19.415	13, 105						
HEMBA1003727   25.803   22.891   29.977   7.994   11.404   10.413   9.955   4.955     HEMBA1003725   10.755   21.218   40.484   11.585   20.280   48.241   48.774   19.547     HEMBA1003726   30.394   76.515   42.218   41.331   22.998   17.475   14.707   20.154   19.547     HEMBA1003737   34.798   43.317   105.267   30.381   19.405   20.814   22.018   22.018   22.855     HEMBA1003737   44.989   43.317   105.267   30.385   19.405   20.814   22.018   22.855     HEMBA1003757   77.780   17.598   12.991   4.750   4.615   7.242   5.982   18.441     HEMBA1003757   77.751   16.033   40.414   11.191   19.441   38.988   28.545   40.914   7.286     HEMBA1003757   77.751   16.033   40.414   11.191   19.441   38.988   28.541   19.288     HEMBA1003757   77.751   16.033   40.414   11.191   19.441   38.988   28.541   19.288     HEMBA1003757   36.779   34.807   113.584   21.315   27.989   17.240   27.704     HEMBA1003757   36.779   34.807   113.584   21.315   27.989   17.240   27.704     HEMBA1003757   36.779   34.807   113.584   21.315   27.989   17.240   27.704     HEMBA1003757   36.779   34.807   113.584   21.315   77.288   20.417   41.485   27.704     HEMBA1003757   36.779   34.807   113.584   21.315   77.288   20.417   41.485   27.704     HEMBA1003768   37.716   36.986   77.222   17.781   17.708   20.417   34.855   26.144   38.986   38.678   37.470   37.478   3		HEMBA1003220	81, 171	86.642	147.453	89.495	42.391	41.580		
HEBBAT003728   105.775   21.728   40.348   11.586   20.280   48.241   44.574   19.547     HEBBAT003720   69.643   70.015   42.439   31.176   20.775   55.815   40.191   27.288     HEBBAT003721   44.989   43.331   105.787   33.038   19.405   20.814   27.018   29.885     HEBBAT003721   44.989   43.331   105.787   33.038   19.405   20.814   27.018   29.885     HEBBAT003725   56.774   57.898   12.939   4.750   4.615   7.242   57.892   4.315     HEBBAT003725   56.774   57.498   12.939   4.750   4.615   7.242   57.892   4.315     HEBBAT003727   71.751   66.033   40.414   11.351   19.441   38.988   28.614   19.028     HEBBAT003727   48.113   38.933   175.742   29.404   21.135   27.2983   17.740   27.704     HEBBAT003727   34.807   31.807   31.812   17.028   20.417   41.885   26.145     HEBBAT003727   31.803   31.522   21.514   10.462   11.887   31.205   15.841   48.55     HEBBAT003273   59.98   24.905   71.222   17.479   15.791   16.787   10.948   17.841     HEBBAT003286   9.746   8.482   12.941   57.79   57.79   37.901   30.931   17.485     HEBBAT003286   9.746   8.482   12.941   57.79   57.79   57.79   57.813   3.552   44.077     HEBBAT003786   9.700   35.057   17.222   17.877   57.79   57.79   57.70   30.931   31.493     HEBBAT003786   9.700   35.057   18.855   37.79   77.79   57.79   57.813   3.555   77.848     HEBBAT003786   9.750   35.931   10.909   3.504   18.100   5.561   5.341   7.547     HEBBAT003786   77.590   5.590   22.443   8.410   17.222   8.448   6.333   9.387     HEBBAT003378   77.590   15.590   22.443   8.410   17.222   8.448   6.333   9.387     HEBBAT003378   77.590   15.590   22.443   8.410   17.222   8.448   6.333   9.387     HEBBAT003378   77.590   15.590   27.443   8.410   17.222   8.448   6.333   9.387     HEBBAT003378   77.590   15.590   27.443   8.410   17.29   8.486   6.331   9.387   17.590     HEBBAT003378   77.590   15.590   27.443   8.410   17.29   8.486   6.331   9.387   17.590     HEBBAT003378   77.590   15.590   27.443   8.410   17.29   8.486   6.331   4.397   17.50     HEBBAT003378					28.577	7,994	11,404	10.413	9.856	14. 985
HEMBA1003225   10.3.1.93								48. 243	44.574	19. 547
HEBBA1003270   65. 641   70.015   42.4319   31.176   20.775   56. 615   40.191   75.238   18.4010   18.4										
HEBBA1003225 49, 499 43, 317 105, 267 33, 308 19, 405 70, 2018 42, 395 8, 11, 165 70, 202 8, 202 18, 241 19, 225 18, 241 19, 226 18, 227 18, 2	15	[HEMBA1003229	30.394							
HEBBA1003226 44,989 43,317 105,267 33,038 19,405 20,814 27,018 29,854 FEBBA1003226 7,760 12,598 12,993 4,750 4,815 7,724 5,255 18,441 FEBBA1003226 7,760 12,598 12,993 4,750 4,815 7,724 5,595 18,441 FEBBA1003226 7,750 15,033 40,414 13,191 19,441 38,988 25,614 19,028 FEBBA1003287 71,751 15,033 40,414 13,191 19,441 38,988 25,614 19,028 FEBBA1003288 19,492 18,398 46,548 41,167 12,756 11,524 6,822 17,414 FEBBA1003276 36,779 34,802 113,584 23,812 17,208 20,417 14,655 25,145 FEBBA1003276 36,779 34,802 113,584 23,812 17,208 20,417 14,655 25,145 FEBBA1003276 36,798 24,906 71,222 17,404 21,135 22,989 17,240 24,704 FEBBA1003276 36,798 24,906 71,222 17,479 15,791 16,787 10,938 17,841 FEBBA1003286 59,507 18,600 38,657 16,933 20,792 37,901 30,931 11,493 FEBBA1003286 59,502 21,393 32,728 15,032 18,415 26,844 28,577 10,348 17,841 FEBBA1003286 59,502 35,947 60,729 21,827 29,473 52,233 50,283 47,695 FEBBA1003286 59,502 35,947 60,729 21,827 29,473 52,233 50,283 47,695 FEBBA1003396 61,539 37,768 48,555 39,127 38,460 40,377 24,057 27,488 FEBBA1003306 77,590 15,530 22,443 8,410 15,787 29,473 52,233 50,283 47,695 FEBBA1003306 77,590 15,530 22,443 8,410 17,786 8,799 34,590 37,768 FEBBA1003306 77,590 15,530 22,443 8,410 17,786 8,799 34,590 37,768 FEBBA1003314 77,117 5,977 8,976 51,530 19,127 38,460 40,377 24,057 27,488 FEBBA1003314 77,117 5,977 8,976 51,530 19,127 38,460 40,377 24,057 27,488 FEBBA1003314 77,117 5,977 8,976 51,530 19,131 19,134 1	13		69, 643	70.015	42.439	31.175	20. 775			
					105, 267	33.038	19, 405	20.834	22.018	29.856
### HEBAN 10037250 7. 260 12. 598 12. 293 4. 150 4. 815 7. 242 5. 982 4. 378									5 295	18. 441
HEIGHA1003255   56.274   51.495   65.197   28.241   33.512   44.917   52.305   60.076		[HEMBA1003250	7.260	12.598!	12.993					
Characteristics   17.75   16.083   40.414   3.391   19.441   38.988   28.614   19.028   19.421   18.988   19.427   18.989   19.421   18.988   19.427   18.989   19.421   18.988   19.427   18.988   19.427   18.988   19.427   18.988   19.427   18.988   19.7240   24.704   19.028   19			56, 274	51, 495	65. 197	28. 241	33.512	44. 917		
							19,441	38, 988	28.614	19.028
HEMBA1003276 36 279 34 802 113 58 4 21 38 52 22 89 17, 220 24, 704 HEMBA1003276 36 279 34 802 113 584 21 812 17 208 20, 417 41, 485 26, 144 455 HEMBA1003278 36 298 74, 906 71, 222 17, 479 15, 791 16, 787 10, 948 17, 841 HEMBA1003280 50, 716 16, 900 38, 607 16, 933 20, 792 37, 790 30, 931 31, 493 11, 494 184 184 184 184 184 184 184 184 184 18	20									
HEIBANO03276 5.6.193 34.802 113.584 23.812 17.208 20.437 14.885 26.145 HEIBANO03277 31.363 17.827 21.514 10.462 11.287 13.206 15.182 14.485 17.841 HEIBANO03280 50.716 16.000 38.057 16.933 20.792 37.901 30.931 31.493 HEIBANO03280 50.716 16.000 38.057 16.933 20.792 37.901 30.931 31.493 HEIBANO03281 65.732 21.393 32.728 15.032 18.415 26.844 28.577 24.898 HEIBANO03286 65.702 21.393 32.728 15.032 18.415 26.844 28.577 24.898 HEIBANO03286 65.002 35.947 60.779 27.827 29.473 52.233 50.238 347.695 HEIBANO03281 69.502 35.947 60.779 27.827 29.473 52.233 50.238 347.695 HEIBANO03294 69.599 52.239 168.555 39.127 38.456 40.377 24.057 27.485 HEIBANO03294 69.599 52.239 168.555 39.127 38.456 40.377 24.057 27.485 HEIBANO03304 7.117 5.972 8.976 6.154 8.839 4.199 3.461 3.227 HEIBANO03304 7.117 5.972 8.976 6.154 8.839 4.199 3.461 3.227 HEIBANO03304 7.117 5.907 28.876 6.154 8.839 4.199 3.461 3.227 HEIBANO03303 6.845 10.103 12.199 14.015 7.776 8.709 3.955 18.326 HEIBANO03303 6.845 10.103 12.199 14.015 7.776 8.709 3.955 18.326 HEIBANO03302 108.401 88.539 256.570 51.038 44.101 1.282 8.448 6.333 9.387 HEIBANO03303 6.845 10.103 12.199 14.015 7.776 8.709 3.955 18.326 HEIBANO03302 108.401 88.539 256.570 51.038 44.101 1.282 8.448 6.333 9.387 HEIBANO03316 37.765 10.128 8.859 37.383 29.387 17.583 17.585 HEIBANO03317 31.501 88.593 256.570 51.038 14.199 7.780 18.802 77.433 17.151 HEIBANO03317 31.501 88.593 256.570 51.038 44.130 48.807 17.450 9.516 HEIBANO03317 315.016 18.7592 87.590 17.780 9.7780 18.699 27.719 7.780 18.699 27.719 7.780 18.699 27.719 7.780 18.699 27.740 9.516 HEIBANO03317 315.016 197.956 389.171 140.044 139.216 140.755 150.455 17.453 18.377 HEIBANO03317 315.016 197.955 389.171 140.044 139.216 140.755 150.455 17.453 18.377 HEIBANO03317 315.016 197.956 389.171 140.044 139.216 140.755 150.455 17.453 18.377 HEIBANO03318 6.568 18.699 19.777 1.126 5.799 36.738 44.891 45.870 17.196 48.117 19.638 17.454 18.4										
HEUBAIO03276 36 279 34.802 113.584 23.812 17.208 20.437 41.805 26.145 HEUBAIO03277 31.355 17.877 24.906 71.222 17.479 15.791 16.787 10.948 17.841 HEUBAIO03278 36.988 24.906 71.222 17.479 15.791 16.787 10.948 17.841 HEUBAIO03280 50.716 16.000 38.057 16.933 20.792 37.901 30.931 31.493 11.246 16.732 21.393 32.728 15.032 18.415 26.844 28.577 24.898 16.804 10.304284 9.746 8.482 12.941 5.779 5.747 5.813 3.545 3.449 16.864 10.3284 9.746 8.482 12.941 5.779 5.747 5.813 3.545 3.449 16.864 10.3284 9.746 8.950 39.947 60.729 21.827 29.473 5.2233 50.283 47.693 17.647 16.864 10.3291 13.248 9.951 10.909 3.504 18.100 6.561 6.341 7.647 16.864 10.3291 13.248 9.951 10.909 3.504 18.100 6.561 6.341 7.647 16.864 16.		HEMBA1003273	48.113	38. 933						
HEBBA1003277 31 363 12.827 21.514 10.462 11.287 13.206 15.182 14.465 HEBBA1003280 50.716 16.000 38.057 16.933 20.792 37.901 30.931 31.493 14.656 16.732 21.393 32.728 15.032 18.415 25.844 28.577 24.838 14.865 16.03284 9.746 8.482 12.941 5.779 5.747 5.813 3.545 3.499 14.8461003284 9.746 8.482 12.941 5.779 5.747 5.813 3.545 3.499 14.8461003286 69.502 35.947 60.729 21.327 29.473 52.233 50.283 47.694 14.8461003291 13.248 9.951 10.909 3.504 18.100 6.551 6.551 6.341 7.647 14.8481003294 69.599 57.239 168.555 39.127 38.460 40.377 24.057 27.488 14.8461003294 69.599 57.239 168.555 39.127 38.460 40.377 24.057 27.485 14.8461003296 61.933 31.456 37.947 21.206 23.199 23.249 34.590 37.768 14.84681003296 61.933 31.456 37.947 21.206 23.199 23.249 34.590 37.768 14.84681003306 7.7.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 14.84681003306 7.7.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 14.84681003316 67.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 14.84681003316 87.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 14.84681003316 87.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 14.84681003316 87.052 210.608 238.618 105.098 198.106 299.884 273.738 171.516 18.84681003315 83.736 51.612 84.690 37.381 72.482 56.594 53.105 54.024 18.8468103315 83.736 51.612 84.690 37.381 72.8462 56.594 53.105 54.024 18.8468103315 83.736 51.612 84.690 37.381 72.8462 56.594 53.105 54.024 18.8468103332 76.8818 107.082 77.78			36, 279	34.802	113.584	23.812	17. 208	20. 437		
#EBBA1003278							11, 287	13. 206	16.182	14.465
#EBBA1003280										
HEMBA10031281   56,7132   21,393   32,728   15,032   18,415   26,844   28,577   24,898   18,406   24,007   24,898   18,406   24,007   24,898   18,406   24,007   24,898   18,406   24,007   24,007   24,057   24,057   24,057   27,486   24,007   27,486   24,007   24,057   27,486   24,007   27,486   27,007   27,486   27,007   27,486   27										
HEMBA1003228		HEMBA1003280	50.716	16.000						
HEMBA1003284 9,746 8,482 12,941 5,779 5,747 5,813 3,545 3,499 HEMBA1003286 69,502 35,947 60,729 21,827 22,947,35,22,333 50,283 47,695 HEMBA1003294 69,599 52,239 168,555 39,127 38,460 40,377 24,057 27,486 HEMBA1003294 69,599 52,239 168,555 39,127 38,460 40,377 24,057 27,486 HEMBA1003304 7,117 5,972 8,976 6,154 8,839 4,199 3,461 3,227 HEMBA1003306 17,590 15,590 22,443 8,410 11,282 8,448 6,333 9,387 168 HEMBA1003314 53,7.052 210,608 238,618 105,008 198,106 299,884 273,738 171,516 HEMBA1003315 83,736 51,812 84,690 32,381 29,482 56,694 53,105 54,024 HEMBA1003315 83,736 51,812 84,690 32,381 29,482 56,694 53,105 54,024 HEMBA1003326 42,723 20,581 14,759 11,799 7,780 18,087 12,420 9,516 HEMBA1003327 61,811 36,702 87,698 28,181 19,780 31,481 30,42 30,44 5,519 HEMBA1003327 61,811 36,702 87,698 28,181 19,784 88,599 22,079 32,865 HEMBA1003327 61,811 36,702 87,698 28,181 19,784 88,599 22,079 32,865 HEMBA1003327 51,811 36,702 87,698 28,181 19,784 88,595 17,453 18,377 HEMBA1003327 51,510 17,114 941 35,926 25,000 18,869 22,079 32,865 HEMBA1003337 315,016 197,955 389,117 14,049 35,926 25,000 18,869 22,079 32,865 HEMBA1003337 35,015 10,275 337,182 94,209 99,716 7,000 43,513 80,023 HEMBA1003376 174,289 170,290 519,668 125,099 89,798 108,226 81,818 107,084 HEMBA1003377 35,015 17,97 53,330 17,430 5,513 19,164 8,117 19,638 HEMBA1003387 42,272 21,380 37,380 42,299 97,17 1,400 544 13,215 14,111 13,485 17,094 14,111 14	25	HEMBA1003281	66.732	21.393	32.728	15.032	18.415	26.844		
HEBBA1003286					12 941	5.779	5,747	5.813	3. 545	3.499
HEMBA1003291 13.248 3.951 10.909 3.504 18.100 6.561 6.341 7.647 HEMBA1003294 69.599 52.239 168.555 39.127 38.460 40.377 24.057 27.486 HEMBA1003204 69.599 52.239 168.555 39.127 38.460 40.377 24.057 27.486 HEMBA1003304 7.117 5.972 8.976 6.154 8.839 4.199 3.4580 37.768 14.060 19.060 1								52 233	50 283	47, 695
HEBBA1003294 69.599 52.239 168.555 39.127 38.460 40.377 24.057 27.486 HEBBA1003296 61.933 31.456 37.947 21.206 23.199 23.249 34.580 37.768 HEBBA1003306 7.117 5.972 8.976 6.154 8.839 4.199 3.461 3.227 HEBBA1003306 7.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 HEBBA1003306 17.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 HEBBA1003306 57.590 15.590 22.443 8.410 11.282 8.448 6.333 9.387 HEBBA1003316 57.592 210.508 23.86 18 105.098 198.106 299.884 273.738 171.516 HEBBA1003315 83.736 51.512 84.690 32.381 29.482 56.594 53.105 54.024 HEBBA1003322 108.401 88.539 256.570 51.502 51.083 44.130 42.804 45.519 HEBBA1003322 42.723 20.581 14.759 17.780 18.087 12.420 9.516 HEBBA1003327 51.811 36.702 87.698 28.181 19.784 18.595 12.409 9.516 HEBBA1003328 53.406 51.712 114.941 35.926 25.000 18.669 22.079 32.865 HEBBA1003328 53.406 51.712 114.941 35.926 25.000 18.669 22.079 32.865 HEBBA1003328 53.806 51.712 114.941 35.926 25.000 18.669 22.079 32.865 HEBBA1003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEBBA10033370 315.016 197.956 369.117 104.044 139.216 140.788 150.458 124.948 HEBBA1003370 315.016 197.956 369.117 104.044 139.216 140.788 150.458 124.948 HEBBA1003370 315.016 197.956 369.117 104.004 139.216 140.788 150.458 124.948 HEBBA1003370 315.016 197.956 369.117 107.004 139.216 140.788 150.458 124.948 HEBBA1003370 11.4269 170.290 519.668 125.099 89.717 67.000 43.513 80.023 HEBBA1003370 315.016 197.956 369.117 107.004 139.216 140.788 150.458 124.948 HEBBA1003370 11.447 25.882 42.659 159.399 89.798 108.228 81.818 107.094 HEBBA1003370 315.016 197.956 369.117 107.004 139.216 140.788 150.458 124.948 HEBBA1003370 315.016 197.956 369.117 107.99 29.81 24.407 19.711 33.485 20.047 HEBBA1003370 315.016 197.956 369.117 107.99 30.5968 129.307 50.088 29.337 23.550 HEBBA1003370 315.016 197.956 369.117 107.99 30.5968 199.596 30.98 108.228 81.818 107.094 140.044 139.30 30.										
HEMBA1003329		HEM8A1003291								
HEMBA1003326   61,933   31,456   37,947   21,206   23,199   23,249   34,580   37,768   16,180A1003306   71,759   15,550   22,443   8,410   11,282   8,448   6,333   9,387   16,180A1003306   17,590   15,550   22,443   8,410   11,282   8,448   6,333   9,387   16,180A1003309   6,845   10,103   12,198   14,015   7,776   8,709   3,955   18,326   16,180A1003314   637,052   210,608   238,618   105,098   198,106   299,884   273,738   171,516   16,180A1003315   83,735   51,612   84,590   32,381   29,482   56,694   53,105   54,024   16,180A1003322   108,401   88,539   256,570   51,502   51,003   44,130   42,804   45,519   16,180A1003327   61,811   36,702   77,598   17,799   7,780   18,087   17,453   18,377   16,180A1003327   61,811   36,702   77,598   28,181   19,784   18,595   17,453   18,377   16,180A1003330   108,955   82,099   207,708   73,413   52,244   50,838   55,920   55,390   16,180A1003369   5,861   23,644   14,930   4,979   1,726   9,064   3,020   5,373   16,180A1003376   315,016   197,956   369,117   140,044   139,216   140,758   150,458   150,458   124,948   16,180A1003373   50,135   31,291   53,330   17,430   4,979   1,726   9,064   3,020   5,373   16,180A1003376   174,269   170,290   519,668   125,099   89,798   108,226   81,818   107,084   16,180A1003376   174,269   170,290   519,668   125,099   89,798   108,226   81,818   107,084   16,180A1003380   43,015   24,657   74,071   29,281   24,407   19,711   13,485   20,047   16,180A1003387   6,515   2,588   2,697   1,577   1,109   1,803   1,986   3,464   1,480A1003399   145,227   21,480   37,035   19,231   24,407   19,711   13,485   20,047   16,180A1003399   45,227   21,480   37,035   19,231   24,557   36,738   44,891   45,870   14,890   44,891   45,870   14,890   44,891   45,870   14,890   44,891   45,870   46,800   46		HEMBA1003294	69, 599	52, 239	168.555	39.127	38.450	40.377		
HEMBA1003304   7.117   5.972   8.976   6.154   8.839   4.199   3.461   3.227     HEMBA1003309   6.845   10.103   12.198   14.015   7.776   8.709   3.955   18.326     HEMBA1003314   537.052   210.608   238.618   105.098   198.106   299.884   273.738   171.516     HEMBA1003315   83.736   51.512   64.690   37.381   29.482   56.694   53.105   54.024     HEMBA1003322   108.401   88.539   256.570   51.502   51.083   44.130   42.804   45.519     HEMBA1003325   42.723   20.581   14.759   11.799   7.780   18.087   12.420   9.516     HEMBA1003328   53.406   51.712   114.941   36.926   25.000   18.669   22.079   32.865     HEMBA1003328   53.406   51.712   114.941   36.926   25.000   18.669   22.079   32.865     HEMBA1003330   108.955   82.099   207.708   73.413   52.244   50.838   55.920   55.390     HEMBA10033370   315.016   197.956   369.117   140.044   139.216   140.758   150.458   124.948     HEMBA1003376   774.265   170.290   519.668   126.099   89.798   108.226   81.818   107.084     HEMBA1003376   774.265   170.290   519.668   126.099   89.798   108.226   81.818   107.084     HEMBA1003387   65.515   25.882   42.253   17.320   29.007   50.086   29.337   23.550     HEMBA1003387   65.515   25.888   42.253   17.320   29.007   50.086   29.337   23.550     HEMBA1003387   65.515   25.882   42.253   17.320   29.007   50.086   29.337   23.550     HEMBA1003395   16.668   18.666   35.483   15.254   9.873   10.355   6.207   12.514     HEMBA1003395   16.068   18.666   35.483   15.254   9.873   10.355   6.207   12.514     HEMBA10033402   37.500   16.239   27.864   8.795   12.867   71.109   1.803   59.453   44.891   44.8				31 456	37,947	21, 206	23. 199	23. 249	34, 580	37. 768
HEMBA1003306 17. 590 15.590 22.443 8.410 11.282 8.448 6.333 9.387   HEMBA1003309 6.845 10.103 12.198 14.015 7.776 8.709 3.555 18.326   HEMBA1003315 83.736 51.612 84.690 32.381 105.098 198.106 299.884 273.738 171.516   HEMBA1003312 108.401 88.539 256.570 51.502 51.083 44.130 42.804 45.519   HEMBA1003327 61.811 36.702 87.698 28.181 19.784 18.596 17.453 18.377   HEMBA1003328 53.406 51.712 114.941 36.926 25.000 18.669 22.079 32.865   HEMBA1003338 53.406 51.712 114.941 36.926 25.000 18.669 22.079 32.865   HEMBA1003338 53.825 82.099 207.708 73.413 52.244 50.838 55.920 55.390   HEMBA1003336 5.861 23.644 14.930 4.979 1.726 9.064 3.020 5.373   HEMBA1003373 50.135 31.291 53.330 17.430 5.513 19.164 8.117 19.638   HEMBA1003373 15.015 197.956 369.117 140.044 139.16 140.758 150.458   HEMBA1003373 50.135 31.291 53.330 17.430 5.513 19.164 8.117 19.638   HEMBA1003373 50.155 31.291 53.330 17.430 5.513 19.164 8.117 19.638   HEMBA1003373 15.015 197.956 369.117 140.044 139.216 140.758 150.458 124.948   HEMBA1003373 50.155 31.291 53.330 17.430 5.513 19.164 8.117 19.638   HEMBA1003376 174.269 170.290 519.668 126.099 89.798 108.226 81.818 107.094   HEMBA1003389 51.6068 18.666 35.483 15.254 99.455 11.810 8.800 14.281   HEMBA1003389 111.457 25.882 42.253 17.323 29.007 50.086 29.337 22.550   HEMBA1003389 111.457 25.882 42.253 17.323 29.007 50.086 29.337 22.550   HEMBA1003389 111.457 25.882 42.253 17.323 29.007 50.086 29.337 22.550   HEMBA1003389 111.457 25.882 42.253 17.323 29.007 50.086 29.337 22.550   HEMBA1003389 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514   HEMBA1003389 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514   HEMBA1003389 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514   HEMBA1003389 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514   HEMBA1003340 116.210 36.907 58.706 24.811 49.133 53.819 60.041 53.109   HEMBA1003402 17.2445 13.970 25.036 8.433 15.354 19.471 27.860 34.116   HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139   HEMBA1003412 104.813 43.934 55.699 77.955 45.336 43.4	20								3.461	3, 227
HEMBA1003309	30									
HEMBA1003314 537.052 210.608 238.618 105.098 198.106 299.884 273.738 171.516   HEMBA1003315 83.736 51.512 84.690 32.381 29.482 56.594 53.105 54.024   HEMBA1003326 188.401 88.539 256.570 51.502 51.083 44.130 42.804 45.519   HEMBA1003327 61.811 36.702 87.698 28.181 19.784 18.596 17.453 18.377   HEMBA1003328 53.406 51.712 114.941 36.926 50.000 18.669 22.079 32.865   HEMBA1003338 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023   HEMBA1003376 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948   HEMBA1003377 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948   HEMBA1003376 174.269 170.290 519.668 126.099 89.718 108.226 81.818 107.084   HEMBA1003384 25.555 30.071 68.079 15.389 9.455 11.810 8.800 14.281   HEMBA1003387 6.515 2.588 2.697 1.577 1.109 1.803 1.986 3.464   HEMBA1003393 11.457 25.382 42.253 17.323 29.007 50.086 29.337 23.550   HEMBA1003393 11.457 25.382 42.253 17.323 29.007 50.086 29.337 23.550   HEMBA1003393 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514   HEMBA1003395 16.688 18.666 35.483 15.254 9.873 10.355 6.207 12.514   HEMBA1003402 37.500 16.219 36.907 58.706 24.811 49.133 55.819 60.041 53.109   HEMBA1003402 37.500 16.239 27.864 8.795 12.867 17.11 11.11		HEMBA1003306	17.590							
HEMBA1003314 637.052 210.608 238.618 105.098 198.106 299.884 273.738 171.516   HEMBA1003312 108.401 88.539 256.597 51.502 51.083 44.130 42.804 45.519   HEMBA1003322 108.401 88.539 256.570 51.502 51.083 44.130 42.804 45.519   HEMBA1003328 42.723 20.581 14.759 11.799 7.780 18.087 12.420 9.516   HEMBA1003328 53.406 51.712 114.941 36.262 25.000 18.669 22.079 32.865   HEMBA1003330 108.955 82.099 207.708 73.413 52.244 50.838 55.920 55.390   HEMBA1003348 121.625 110.275 337.182 94.209 99.117 67.000 43.513 80.023   HEMBA1003370 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948   HEMBA1003370 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948   HEMBA1003376 174.269 170.290 519.688 126.099 8.798 108.226 81.818 107.084   HEMBA1003380 43.015 24.657 74.071 29.281 24.407 19.711 13.485 20.047   HEMBA1003387 50.155 30.071 68.079 15.389 9.455 11.810 8.800 14.281   HEMBA1003387 50.515 2.588 2.697 1.577 1.109 1.803 1.986 3.464   HEMBA1003389 45.227 21.480 37.035 19.231 15.354 19.171 17.355 6.207 12.514   HEMBA1003399 45.227 21.480 37.035 19.231 15.354 19.471 27.860 34.116   HEMBA1003340 37.500 16.239 27.864 8.795 12.867 17.141 1.617 14.595   HEMBA1003402 37.500 16.239 27.864 8.795 12.867 17.141 1.617 1.4596   HEMBA1003403 16.210 36.907 58.706 24.811 49.133 53.819 60.041 53.109   HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.385 77.062 46.433   HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.385 77.062 46.433   HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.385 77.062 46.433   HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.385 77.062 46.433   HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.385 77.062 46.433   HEMBA1003407 17.22.445 13.970 25.036 8.433 7.282 10.593 5.593 5.696 11.032   HEMBA1003407 17.466 15.895 21.662 4.733 6.721 8.483 10.838 9.083   HEMBA1003412 17.466 15.895 21.662 4.733 6.721 8.483 10.838 9.083   HEMBA1003433 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581		HEMBA1003309	6.845	10, 103	12.198	14.015				
HEMBA1003315 83.736 51.612 84.690 32.381 29.482 56.694 53.105 54.024 HEMBA1003326 108.401 88.539 255.570 51.502 51.083 44.130 42.804 45.519 HEMBA1003327 61.811 36.702 87.698 28.181 19.784 18.596 17.453 18.377 HEMBA1003328 53.406 51.712 114.941 36.926 25.000 18.669 22.079 32.865 HEMBA1003330 108.955 82.099 207.708 73.413 52.244 50.838 55.920 55.390 HEMBA1003330 108.955 82.099 207.708 73.413 52.244 50.838 55.920 55.390 HEMBA1003369 5.861 23.644 14.930 4.979 1.726 9.064 3.020 5.373 HEMBA1003370 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948 HEMBA1003376 174.269 170.290 519.668 125.099 89.717 67.000 43.513 80.023 HEMBA1003376 174.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003376 174.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003376 574.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003376 574.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003376 574.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003376 574.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003378 50.136 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBA1003395 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBA1003395 15.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBA1003308 195.676 49.687 70.460 29.354 9.873 10.355 6.207 12.514 HEMBA1003402 12.2500 16.239 27.864 8.795 12.867 17.141 11.617 14.595 HEMBA1003402 12.2500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 12.2500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 12.2500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 12.2500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 12.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 12.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.593 59.463 47.139 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMB				210,608	238,618	105,098	198, 106	299.884	273.738	171, 516
HEMBA1003322 108. 401 88. 539 255. 570 51. 502 51. 083 44. 130 42. 804 45. 519 HEMBA1003326 42. 723 20. 581 14. 759 11. 799 7. 780 18. 087 12. 420 9. 516 HEMBA1003327 61. 811 36. 702 87. 698 28. 181 19. 784 18. 596 17. 453 18. 377 HEMBA1003328 53. 406 51. 712 114. 941 36. 926 25. 000 18. 669 22. 079 32. 865 HEMBA1003330 108. 955 82. 099 207. 708 73. 413 52. 244 50. 838 55. 920 55. 390 HEMBA1003348 121. 625 110. 275 337. 182 94. 209 99. 717 67. 000 43. 513 80. 023 HEMBA1003370 315. 016 197. 956 369. 117 140. 044 139. 216 140. 758 150. 458 124. 948 HEMBA1003373 50. 135 31. 291 53. 330 17. 430 5. 513 19. 164 8. 117 19. 638 HEMBA1003376 174. 269 170. 290 519. 668 126. 099 89. 798 108. 226 81. 818 107. 084 HEMBA1003376 174. 269 170. 290 519. 668 126. 099 89. 798 108. 226 81. 818 107. 084 HEMBA1003376 6. 515 2. 588 2. 697 15. 389 9. 455 11. 810 8. 800 14. 281 HEMBA1003378 6. 515 2. 588 2. 697 1. 577 1. 109 1. 803 1. 986 3. 464 HEMBA1003395 16. 668 18. 666 35. 463 15. 254 9. 873 10. 355 6. 207 12. 516 HEMBA1003395 45. 227 21. 480 37.035 19. 231 15. 354 19. 471 27. 860 34. 116 HEMBA1003400 116. 210 36. 907 58. 706 24. 811 49. 133 53. 819 60. 041 53. 109 HEMBA1003402 32. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 HEMBA1003402 37. 500 16. 239 27. 864 8. 795 12. 867 17. 141 11. 617 14. 596 14. 800 14. 80							29 482	56, 694	53, 105	54.024
HEMBA1003326   42, 723   20, 581   14,759   1,799   7,780   18,087   12,420   9,516     HEMBA1003327   61,811   36,702   87,658   28,181   19,784   18,596   17,453   18,377     HEMBA1003328   53,406   51,712   114,941   36,926   25,000   18,669   22,079   32,865     HEMBA1003330   108,955   82,099   207,708   73,413   52,244   50,838   55,920   55,390     HEMBA1003348   121,625   110,275   337,182   94,209   99,717   67,000   43,513   80,023     HEMBA1003370   315,016   197,956   369,117   140,044   139,216   140,758   150,458   124,948     HEMBA1003373   50,135   31,291   53,330   17,430   5,513   19,164   8,117   19,638     HEMBA1003376   174,269   170,290   519,668   126,099   89,798   108,226   81,818   107,084     HEMBA1003384   25,555   30,071   68,079   15,389   9,455   11,810   8,800   14,281     HEMBA1003395   111,457   25,382   42,253   17,323   29,007   50,086   29,337   23,550     HEMBA1003395   16,668   18,666   35,483   15,254   9,873   10,355   6,207   12,514     HEMBA1003395   16,668   18,666   35,483   15,254   9,873   10,355   6,207   12,514     HEMBA1003340   116,210   36,907   58,706   48,811   49,133   53,819   60,041   53,109     HEMBA1003400   116,210   36,907   58,706   48,811   49,133   53,819   60,041   53,109     HEMBA1003403   60,260   43,377   46,720   20,221   26,579   36,738   44,891   45,870     HEMBA1003403   106,813   43,934   55,699   47,250   43,763   51,953   59,463   47,139     HEMBA1003402   27,544   33,970   25,036   8,433   7,282   10,593   5,696   11,032     HEMBA1003412   104,813   43,934   55,699   47,250   43,763   51,953   59,463   47,139     HEMBA1003412   104,813   43,934   55,699   47,250   43,763   51,953   59,463   47,139     HEMBA1003412   104,813   43,934   55,699   47,250   43,763   51,953   59,463   47,139     HEMBA1003412   104,813   43,934   55,699   47,250   43,763   51,953   59,463   47,139     HEMBA1003418   57,411   57,397   76,232   97,795   45,336   43,450   22,206   90,604     HEMBA1003425   17,466   15,895   21,662   4,733   6,723   8,483   10,3										
HEMBA1003327 61.811 36.702 87.698 28.181 19.784 18.596 17.453 18.377 HEMBA1003328 53.406 51.712 114.941 36.926 25.000 18.669 22.079 32.865 HEMBA1003338 108.955 82.099 207.708 73.413 52.244 50.838 55.920 55.390 HEMBA1003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEMBA1003370 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948 HEMBA1003375 50.135 31.291 53.330 17.430 5.513 19.164 8.117 19.638 HEMBA1003376 174.269 170.290 519.668 125.099 89.798 108.226 81.818 107.084 HEMBA1003384 25.555 30.071 68.079 15.389 9.455 11.810 8.800 14.281 HEMBA1003387 6.515 2.588 2.697 1.577 1.109 3.803 1.986 3.464 HEMBA1003395 16.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBA1003399 45.227 21.480 37.035 19.231 15.354 19.471 27.860 34.116 HEMBA1003309 45.227 21.480 37.035 19.231 15.354 19.471 27.860 34.116 HEMBA1003402 32.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.431 HEMBA1003408 196.676 49.687 70.460 19.807 11.379 12.938 14.721 HEMBA1003408 196.676 49.687 70.										
HEMBA1003327 61.811 36.702 87.698 28.181 19.784 18.595 17.433 18.377 HEMBA1003328 53.406 51.712 114.941 36.926 25.000 18.669 22.079 32.865 HEMBA1003330 108.955 82.099 207.708 73.413 52.244 50.838 55.920 55.390 HEMBA1003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEMBA1003369 5.861 23.644 14.930 4.979 1.726 9.064 3.020 5.373 HEMBA1003370 315.015 197.956 369.117 140.044 139.216 140.758 150.458 124.948 HEMBA1003373 50.135 31.291 53.330 17.430 5.513 19.164 8.117 19.638 HEMBA1003376 174.269 170.290 519.668 125.099 89.795 108.226 81.818 107.084 HEMBA1003380 43.015 24.657 74.071 29.281 24.407 19.711 13.485 20.047 HEMBA1003387 6.515 2.588 2.697 1.577 1.109 1.803 1.986 3.464 HEMBA1003392 111.457 25.882 42.253 17.323 29.007 50.086 29.337 23.550 HEMBA1003399 45.227 21.480 37.035 19.231 15.354 19.471 27.860 34.116 HEMBA1003402 116.210 36.907 58.706 24.811 49.133 53.819 60.041 53.109 HEMBA1003402 37.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003403 195.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 51.953 59.463 47.139 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 51.953 59.463 47.139 HEMBA1003402 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003405 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003405 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003405 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003405 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003405 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083	35	HEMBA1003326	42.723	20.581	14.759					
HEMBAT003328 53.406 51.712 114.941 36.926 25.000 18.669 22.079 32.865 HEMBAT003330 108.955 82.099 207.708 73.413 52.244 50.838 55.920 55.390 HEMBAT003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEMBAT003369 5.861 23.644 14.930 4.979 1.726 9.064 3.020 5.373 HEMBAT003370 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948 HEMBAT003376 174.269 170.290 519.668 126.099 89.798 108.226 81.818 107.084 HEMBAT003380 43.015 24.657 74.071 29.281 24.407 19.711 13.485 20.047 HEMBAT003387 6.515 24.687 74.071 29.281 24.407 19.711 13.485 20.047 HEMBAT003387 6.515 2.588 2.697 1.577 1.109 1.803 1.986 3.464 HEMBAT003395 16.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBAT003395 16.068 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBAT003340 45.277 21.880 37.035 19.231 15.354 19.471 27.860 34.116 HEMBAT003340 116.210 36.907 58.706 24.811 49.133 53.819 60.041 53.109 HEMBAT003402 32.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBAT003403 196.676 49.687 70.460 29.354 50.910 84.358 77.062 44.891 HEMBAT003417 22.445 13.970 25.036 8.433 7.282 10.593 59.463 47.139 HEMBAT003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBAT003425 17.466 15.895 2		HEMBA1003327	61.811	36, 702	87.698	28. 181	19, 784	18.596		
HEMBA1003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEMBA1003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEMBA1003369 5.861 23.644 14.930 4.979 1.726 9.064 3.020 5.373 HEMBA1003370 315.016 197.956 369.117 140.044 139.216 140.758 150.458 124.948 HEMBA1003373 50.135 31.291 53.330 17.430 5.513 19.164 8.117 19.638 HEMBA1003376 174.269 170.290 519.668 126.099 89.798 108.226 81.818 107.084 HEMBA1003380 43.015 24.657 74.071 29.281 24.407 19.711 13.485 20.047 HEMBA1003387 6.515 2.588 2.697 15.389 9.455 11.810 8.800 14.281 HEMBA1003392 111.457 25.882 42.253 17.323 29.007 50.086 29.337 23.550 HEMBA1003395 16.668 18.666 35.483 15.254 9.873 10.355 6.207 12.514 HEMBA1003399 45.227 21.480 37.035 19.231 15.354 19.471 27.860 34.116 HEMBA1003402 115.210 36.907 58.706 24.811 49.133 53.819 60.041 53.109 HEMBA1003403 50.260 43.377 46.720 20.221 26.579 36.738 44.891 45.870 HEMBA1003403 195.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003403 150.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003403 150.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003403 150.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003403 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003402 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003402 17.466 15.885 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.885 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.885 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.885 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA10034325 17.466 15.885 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA10034325 17.466 15.885 21.662 4.733 6.723						36 926	25,000	18.669	22.079	32.865
HEMBA1003348 121.625 110.275 337.182 94.209 99.717 67.000 43.513 80.023 HEMBA1003369 5.861 23.644 14.930 4.979 1.726 9.064 3.020 5.373 HEMBA1003370 315.016 197.956 369.117 140.044 139.216 140.768 150.458 124.948 HEMBA1003376 174.269 170.290 519.668 126.099 89.783 19.164 8.117 19.638 HEMBA1003376 174.269 170.290 519.668 126.099 89.798 108.226 81.818 107.084 HEMBA1003380 43.015 24.657 74.071 29.281 24.407 19.711 13.485 20.047 HEMBA1003387 6.515 2.588 2.697 1.577 1.109 1.803 1.986 3.464 HEMBA1003392 111.457 25.882 42.253 17.323 29.007 50.086 29.337 23.550 HEMBA1003392 150.668 18.666 35.463 15.254 9.873 10.355 6.207 12.514 HEMBA1003395 45.227 21.480 37.035 19.231 15.354 19.471 27.860 34.116 HEMBA1003400 116.210 36.907 58.706 24.811 49.133 53.819 60.041 53.099 HEMBA1003402 32.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003403 196.676 49.887 70.460 29.354 50.910 84.358 77.062 45.433 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.9463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.								50 838	55, 920	55, 390
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HEMBA1003402 32.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003402 32.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003403 60.260 43.277 46.720 20.221 26.579 36.738 44.891 45.870 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083		HEM8A1003399	45, 227	21,480	37.035	19.231	15. 354	19.471	27.860	
HEMBA1003402 32.500 16.239 27.864 8.795 12.867 17.141 11.617 14.596 HEMBA1003403 60.260 43.377 46.720 20.221 26.579 36.738 44.891 45.870 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 55.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.855 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003423 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581							49, 133	53.819	60.041	53. 109
HEMBA1003402 35.00 43.377 46.720 20.221 26.579 36.738 44.891 45.870 HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003433 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581										14 596
HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 46.433 HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003433 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581										
HEMBA1003408 196.676 49.687 70.460 29.354 50.910 84.358 77.062 45.433 HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 55.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003433 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581		HENBA 1003403	60.260	1 43.377	45.720					
HEMBA1003412 104.813 43.934 55.699 47.250 43.763 61.953 59.463 47.139 HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003433 23.931 18.415 24.576 12.136 10.421 10.074 11.092 11.581				49.687	70.460	29.354	50.910	84.358		
HEMBA1003417 22.445 13.970 25.036 8.433 7.282 10.593 5.696 11.032 HEMBA1003418 57.411 57.397 76.232 97.795 45.336 43.450 22.206 90.604 HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003433 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581	50							61.953	59, 463	47. 139
HEMBA1003418 57, 411 57, 397 76, 232 97, 795 45, 336 43, 450 22, 206 90, 604 HEMBA1003420 29, 838 15, 856 201, 831 11, 319 8, 067 11, 379 12, 938 14, 721 HEMBA1003425 17, 466 15, 895 21, 662 4, 733 6, 723 8, 483 10, 838 9, 083 HEMBA1003433 23, 931 18, 435 24, 576 12, 136 10, 421 10, 074 11, 092 11, 581										
HEMBA1003420 29.838 15.856 201.831 11.319 8.067 11.379 12.938 14.721 HEMBA1003425 17.466 15.895 21.662 4.733 6.723 8.483 10.838 9.083 HEMBA1003433 23.931 18.435 24.576 12.136 10.421 10.074 11.092 11.581										
HEMBA1003425 17, 466 15,895 21,662 4,733 6,723 8,483 10,838 9,083 HEMBA1003433 23,931 18,435 24,576 12,136 10,421 10,074 11,092 11,581		HEMBA1003418	57, 411	57.397	76.232	97.795				
HEMBA1003425 17, 466 15, 895 21, 662 4, 733 6, 723 8, 483 10, 838 9, 083 HEMBA1003433 23, 931 18, 435 24, 576 12, 136 10, 421 10, 074 11, 092 11, 581				15.856	201.831	11.319	8.057	11.379	12.938	
HEMBA1003433 23, 931 18, 415 24, 576 12, 136 10, 421 10, 074 11, 092 11, 581								8.483	10.838	9. 083
[[EMDATOD433   Ex. 331   10. 433   Ex. 434   E										
35	5.5	HEMBA1003433	23.931	18.43	24.5/	12.130	1 10.421	1 10.014	1 11. 032	
	JJ									

Table 26

	THE PART OF T	A) 747 [	41 727 1	39.257	19.755	26, 941	45, 998	31.620	35.845
	HENBA1003440	91. 727	41.727	10.452	33.897	10. 259	15. 118	7.093	14.790
	HEMBA1003442	7.090	22. 535	48. 248	26.789	18. 587	41. 591	42.314	35.065
5	HEMBA1003447	82. 151	36.670	25. 954	16. 130	11, 252	16. 584	28. 534	21.256
J	HEMBA1003453	50. 472	26.692	42.686	17. 261	18.856	27. 281	22.795	17.854
	HEMBA1003461	55. 687	25. 328	34. 469	13.456	19.704	20. 277	16.984	18.124
	HEMBA1003463	40.102	23.311	51.816	28. 410	36.051	39, 389	40. 220	36. 851
	HEMBA1003465	92. 245	40.963	266.076	76, 166	67.942	56. 459	51.589	62. 191
	HEMBA1003480	114.075	114.841	33.659	14. 371	8. 636	26. 284	16.036	14. 582
10	HEMBA 1003485	44, 403	28.836		9, 752	15.729	24. 902	21.136	16.494
10	HENBA 1003487	42.939	15.463	23.730 56.674	14. 934	12.014	12.082	9. 567	14.655
	HEMBA1003492	31.026	21.538		48. 821	12.504	74. 554		180, 841
	HEMBA 1003494	97.366	260.496	50.174 24.659	11, 432	13.881	21.376	18.562	8.072
	HEMBA1003497	39.000	17.943	28. 175	12. 948	17, 154	30.911	36.453	16.806
	HEMBA1003503	54.774 18.672	21.486	43.023	11.794	13. 330	8. 925	16.405	11.615
15	HEMBA1003511		191.234	239.319	81.329	123.915	213.945	179.430	96. 672
,,	HEMBA 1003528	385.123 43.820	12.384	23.693	10,695	21.216	20.067	28.030	16. 204
	HEMBA 1003530 HEMBA 1003531	111, 104	73.542	215.578	67.833	214.022	56.139	50. 217	66.992
	HENBA1003532	145. 137	62. 379,	83.827	37.506	53.388	90.314	77.728	60. 515
	HENBA1003538	61. 123	20.746	32,949	11.180	19. 286	34. 305	28. 231	13. 837
	HEMBA1003545	21.489	10.501	20,608	5. 904	7.197	10.239	6.617	8. 168
20	HEMBA1003546	31.371	32.365	28.613	13.365	226. 243	16.427	16.554	24.821
_•	HEMBA1003548	4, 466	8. 124	9.845	4. 563	7.542	6.155	5. 647	8.387
	HEMBA1003553	79.837	51.515	50.379	23. 327	28.564	49. 154	63. 525	48. 955
	HEMBA1003555	20.066	8.873	13.692	4.762	3.684	10.112	10.962	6. 521
	HEMBA1003556	57. 280	36.399	128.391	29. 283	16.426	19. 257	18. 121	24.622
	HEMBA1003560	9. 290	4.426	2.529	2.848	1,767	2.983	6. 207	6.539
25	HEMBA1003565	42.648	29.588	20.996	8, 344	13.984	21.927	21.847	22.043
	HEMBA1003568	7.244	1.649	7.712	2. 430	3.763	3. 172	2.836	2. 592 16. 518
	HEMBA1003569	25.048	20.536	23.764	33. 957	13.740	16. 235	19.512 48.011	59. 960
	HEMBA 1003571	111.721	94. 378	326.335	84. 368	71.788	50.029 2.872	9. 198	6. 421
	HEMBA1003579	3. 335	7.399	15.353	6, 553	59.875	128.943	110.375	35. 695
	HENBA1003580	274. 105	50. 292	102.103	26.686	36.215	54, 336	50.711	21.238
30	HEMBA 1003581	112.013	31. 295	94.083	21.641 89.876	47.882	53.615	40, 656	45. 172
	HEMBA 1003591	97.076	64. 326	84.629	19.075	11.339	6.305	5. 581	18.085
	HENBA1003595	32.697	22.842 25.846	108.491	20.931	15. 952	19.375	17.580	20. 153
	HEMBA1003597	48.561	20. 134	22.468	12.142	11.688	18.934	21.743	15.025
	HEMBA1003598 HEMBA1003600	49.728 32.772	35.099	56. 905	26.268	29.290	38.873	53. 305	56.783
	HENBA1003602	18. 248	10.116	16, 162	6. 182	10.970	8.064	14.736	17. 188
35	HEMBA1003604	205. 949	53. 579	69,723	24. 549	49. 902	105.181	98. 166	47. 144
	HEMBA1003610	140. 996	29. 255	95.048	15. 492	103.150	72.233	54.670	30.688
	HEMBA1003615	57.258	20.035	34. 102	12.808	16.022	24. 378	18.759	20.876
	HEMBA1003617	48.414	20.375	29,789	12.148	22.291	18.199	18.770	18. 242
	HEMBA1003620	52.899	22.318	45. 502	19.575	19.962	25.239	39.072	29. 451
	HEMBA1003621	102.827	102.094	226, 373	80.194	64.742	58.874	67.142	60.680
40	HEMBA1003622	19.815	13.838	25.009	16.055	8. 339	12.261	15. 369	13.833
	HEMBA1003630	20.008	16.381	30.244	13.871	5. 573	9, 992	10.303	18. 500
	HEMBA1003637	37.880	29.848		23. 251	18.468	17. 181	17.417	20. 420
	HEMBA 1003640	39.068	31.672		22.572	22.223	21.513	54. 230	5,711
	HEMBA1003645	25,820	19.380			9. 247	24.606	19. 177	19.938
	HEM8A 1003646	38. 243	16. 329				7.507	7. 882	10.058
45	HEMBA1003647	10.261	10.718					23. 172	30, 178
	HEMBA 1003656	40, 171					10.909		18. 544
	HEMBA1003662	25. 325					9.774		13.851
	HEMBA1003666	23.086					179.317	140.769	174. 256
	HEMBA1003667	304.975					7.057		
50	HEMBA 1003670	12.944					66,675		
50	HEMBA 1003674	80.516							
	HEMBA 1003677 HEMBA 1003679	25. 325							
	HEMBA1003680	42.317							
	HEMBA1003684	18. 273						10. 182	
	HEMBA1003690							85.431	56.592

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Table 27

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HEMBA1003692	83.253	96. 347   1	94.372	63. 188	40.872	47.354	39.288	42.644
HEMBA1003702	88. 125	35.028		23.719	29.023	42.879	45.956	36.550
				32. 886	33. 424	47.500	41, 959	36.807
HEMBA1003711	93.732	****			28. 237	32, 029	29.145	16.214
HEMBA1003714	75. 923	20.696		14. 414				
HEMBA1003715	54, 160	54.486	42.871	31.894	31.122	28. 832	20.640	26.672
HEMBA1003717	70, 553	38. 574	20.922	45, 101	29.491	29. 344	27. 200	38.418
	83. 687			55. 896	49.519	43, 330	22.099	41, 137
HEMBA1003720			71.704	30.085	21. 305	22.378	18.643	31.573
HEMBA1003725	46.157	55. 932				42. 286	37.790	34. 280
HEMBA1003728	103.795	35. 668	58. 184	16.485	21.818			
HEMBA1003729	49, 957	21.508	47.663	20. 231	15. 376	18. 567	21. 294	17. 427
HEMBA1003732	13.069	1.953	6.558	3. 228	2. 195	3.652	3.024	4. 336
HEMBA1003733	52. 409	32.781	76.684	22.919	83.426	18.921	13.867	14. 220
			50.667	26. 589	21.518	42.057	44.130	24.802
HEMBA1003742	40.426	20. 265				12.506	17.162	18.069
HEMBA1003743	26.918	22.118	23. 392	18.886	18.530			
HEMBA1003758	110,630	126.359	315.104	79. 435	58. 130	58. 587	34.868	73.429
HEMBA1003760	78.949	0.000	26.318	15, 194	14.440	32.057	34. 468	19.471
		30, 390	82.720	23.891	19.630	164, 051	37.797	57.861
HEMBA1003764	45.855				32.047	46, 499	39.296	38.944
HEMBA1003769	87.589	47.227	62.942	27.144				
HEMBA1003773	63.842	14.722	21.132	12.002	9.850	33. 904	29.817	13. 165
HEMBA1003783	17,751	16. 975	23.942	16.465	13.884	6.842	9.757	20.650
HEMBA1003784	13.500	17. 233	21.849	13, 856	12.436	17.394	11.099	13.140
		303.008		109.371	145.316	286.778	287.377	239.938
HEMBA1003794	386.642			15.022	13.775	16. 550	24.428	19. 403
HEMBA1003799	39, 392	23.099	29.603					35. 938
HEMBA 1003803	63.548	21.899	44. 323	20.132	18.580	28. 795	24.744	
HEMBA1003804	80.382	26.816	48, 558	15.154	27.867	31.087	37.611	22.634
HEMBA1003805	103.669	42, 485	42, 930	19.994	36.377	43.797	32.147	28.376
	21.717	13.940	25, 512	9.492	6.870	9. 649	8,812	7.611
HEMBA 1003807				20. 338	17.855	7.640	4, 451	6. 585
HEMBA1003810	20, 102	11.572	7. 558					
HEMBA 1003827	432.964	219.520		155.416	219.584	266.037	283, 204	241. 127
HE LEBA 1003836	177.311	135.831	482.334	146.466	136.063	93.790	92.728	122. 237
HEMBA1003838	223.674	185. 295	641.368	134.002	79.993	115.711	87.137	118.957
HEMBA1003843	13.867	10. 178	27. 409	17.850	21.104	13.382	11.701	13.634
				34, 962	50.550	56. 395	40.851	60. 253
HEMBA1003846	133.994	57. 556	58.738			11, 259		9.934
HEMBA1003856	27.378	13.868	16.982	14.248	8.662		9.145	
HEMBA1003857	101,908	95, 527	253, 525	75. 110	52.628	51.958	45.837	48.871
HEMBA1003864	52, 130	18.071	24.567	9, 568	13.009	16.810	29.271	16.795
HEMBA1003866	27.257	12.805	22.440	12.069	15.414	19, 103	9.229	7.524
			58. 923	31.090	41.733	69.461	48.174	43.486
HEMBA1003868	95.701	54.991				37.097	36.010	45.824
HEMBA1003879	62.950	44. 572	159. 217	48.098	42.446		44, 498	30.581
HEMBA1003880	134.462	70, 074	103.271	50,699	47.956	67.668		
HEMBA1003884	99.190	48. 465	73.499	34.796	54.399	57.269	63.551	68.830
HEMBA1003885	77.675	69.096	172.968	55, 129	49.424	41.309	24. 247	31.596
HEMBA1003887	60. 203	22. 185	33.582	16.896	21.181	29. 281	31.275	22.835
		8. 056	15, 506	7.762	16.057	139.271	387.408	5.124
HEMBA1003890	12.753				180.355	212.964	137.297	122, 335
HEMBA1003893	386.525	281.955	515.307	187.300			185.700	143.577
HEMBA1003896	411.418	232.899	382. 182	144, 104	165. 806	233.857		
HEMBA1003902	39.732	39.491	114.984	20.297	23.509	16.793	14.124	20.479
HENBA1003904	32.775	21.109	45.629	10.006	13.109	14. 294	24. 342	17.444
HEMBA 1003908	8. 560	8.873	15.689	7.298	15. 429	6.307	2.267	5.699
		253.614		183.017	124.195	147, 955	105.962	360.995
HEMBA 1003926	132.636		316.882			35. 284	29.595	40, 418
HEMBA 1003937	87.005	63.862	200. 940	40.687	36.238			15, 626
	1 AA AF A	1 25 044	35.675	20.306	20.378	19.070	16.457	
HEMBA1003939	28.064	25.844						9.868
HEMBA1003940	27,800	13.368	18.045	10.235	10.394	14.633	17.733	
HEMBA1003940	27,800		18.045	10.235		14. 633 23. 428		13.795
HEMBA 1003940 HEMBA 1003941	27.800 57.997	13, 368 16, 835	18. 045 24. 582	17.381	15.884	23. 428		
HEMBA 1003940 HEMBA 1003941 HEMBA 1003942	27.800 57.997 38.168	13.368 16.835 19.747	18.045 24.582 45.852	17.381 32.660	15.884 22.333	23. 428 24. 695	19.757 10.791	13.795 21.900
HEMBA 1003940 HEMBA 1003941 HEMBA 1003942 HEMBA 1003945	27.800 57.997 38.168 59.457	13.368 16.835 19.747 32.900	18.045 24.582 45.852 46.079	17. 381 32. 660 23. 037	15.884 22.333 21.163	23. 428 24. 695 36. 632	19.757 10.791 32.279	13.795 21.900 26.903
HEMBA 1003940 HEMBA 1003941 HEMBA 1003942	27.800 57.997 38.168	13. 368 16. 835 19. 747 32. 900 13. 019	18.045 24.582 45.852 46.079 20.678	17.381 32.660 23.037 7.159	15.884 22.333 21.163 38.521	23. 428 24. 695 36. 632 442. 120	19.757 10.791 32.279 272.494	13.795 21.900 26.903 21.625
HEMBA 1003940 HEMBA 1003941 HEMBA 1003942 HEMBA 1003945	27.800 57.997 38.168 59.457	13.368 16.835 19.747 32.900	18. 045 24. 582 45. 852 46. 079 20. 678 5. 814	17.381 32.660 23.037 7.159 3.195	15.884 22.333 21.163 38.521 4.756	23. 428 24. 695 36. 632 442. 120 3. 396	19.757 10.791 32.279 272.494 8.814	13.795 21.900 26.903 21.625 5.401
HEMBA 1003940 HEMBA 1003941 HEMBA 1003942 HEMBA 1003945 HEMBA 1003949 HEMBA 1003950	27.800 57.997 38.168 59.457 12.870 8.366	13. 368 16. 835 19. 747 32. 900 13. 019 8. 726	18. 045 24. 582 45. 852 46. 079 20. 678 5. 814	17.381 32.660 23.037 7.159 3.195	15.884 22.333 21.163 38.521	23. 428 24. 695 36. 632 442. 120	19.757 10.791 32.279 272.494 8.814	13.795 21.900 26.903 21.625
HEMBA 1003940 HEMBA 1003941 HEMBA 1003942 HEMBA 1003945 HEMBA 1003949 HEMBA 1003950 HEMBA 1003953	27.800 57.997 38.168 59.457 12.870 8.366 23.527	13. 168 16. 835 19. 747 32. 900 13. 019 8. 726 10. 310	18.045 24.582 45.852 46.079 20.678 5.814 11.872	17.381 32.660 23.037 7.159 3.195 9.390	15.884 22.333 21.163 38.521 4.756 8.494	23. 428 24. 695 36. 632 442. 120 3. 396	19.757 10.791 32.279 272.494 8.814 10.973	13.795 21.900 26.903 21.625 5.401
HEMBA1003940 HEMBA1003941 HEMBA1003942 HEMBA1003945 HEMBA1003950 HEMBA1003950 HEMBA1003953 HEMBA1003958	27.800 57.997 38.168 59.457 12.870 8.366 23.527 131.082	13, 168 16, 835 19, 747 32, 900 13, 019 8, 726 10, 310 90, 718	18.045 24.582 45.852 46.079 20.678 5.814 11.872 253.084	17.381 32.660 23.037 7.159 3.195 9.390 74.499	15.884 22.333 21.163 38.521 4.756 8.494 85.036	23. 428 24. 695 36. 632 442. 120 3. 396 10. 637 52. 450	19.757 10.791 32.279 272.494 8.814 10.973 34.852	13.795 21.900 26.903 21.625 5.401 5.252 86.629
HEMBA1003940 HEMBA1003941 HEMBA1003942 HEMBA1003945 HEMBA1003954 HEMBA1003953 HEMBA1003958 HEMBA1003958	27.800 57.997 38.168 59.457 12.870 8.366 23.527 131.082 12.105	13.368 16.835 19.747 32.900 13.019 8.726 10.310 90.718	18. 045 24. 582 45. 852 46. 079 20. 678 5. 814 11. 872 253. 084 18. 520	17. 381 32. 660 23. 037 7. 159 3. 195 9. 390 74. 499 6. 548	15.884 22.333 21.163 38.521 4.756 8.494 85.036 7.960	23. 428 24. 695 36. 632 442. 120 3. 396 10. 637 52. 450 18. 122	19.757 10.791 32.279 272.494 8.814 10.973 34.852 12.612	13.795 21.900 26.903 21.625 5.401 5.252 86.629 10.591
HEMBA1003940 HEMBA1003941 HEMBA1003942 HEMBA1003945 HEMBA1003945 HEMBA1003950 HEMBA1003953 HEMBA1003959 HEMBA1003959	27.800 57.997 38.168 59.457 12.870 8.366 23.527 131.082 12.105 53.133	13, 368 16, 835 19, 747 32, 900 13, 019 8, 726 10, 310 90, 718 11, 228 29, 785	18. 045 24. 582 45. 852 46. 079 20. 678 5. 814 11. 872 253. 084 18. 520 31. 879	17. 381 32. 660 23. 037 7. 159 3. 195 9. 390 74. 499 6. 548 18. 932	15.884 22.333 21.163 38.521 4.756 8.494 85.036 7.960 16.178	23. 428 24. 695 36. 632 442. 120 3. 396 10. 637 62. 450 18. 122 21. 708	19.757 10.791 32.279 272.494 8.814 10.973 34.852 12.612 32.094	13.795 21.900 26.903 21.625 5.401 5.252 86.629 10.591 35.333
HEMBA1003940 HEMBA1003941 HEMBA1003942 HEMBA1003945 HEMBA1003954 HEMBA1003953 HEMBA1003958 HEMBA1003958	27.800 57.997 38.168 59.457 12.870 8.366 23.527 131.082 12.105	13.368 16.835 19.747 32.900 13.019 8.726 10.310 90.718	18. 045 24. 582 45. 852 46. 079 20. 678 5. 814 11. 872 253. 084 18. 520	17. 381 32. 660 23. 037 7. 159 3. 195 9. 390 74. 499 6. 548	15. 884 22. 333 21. 163 38. 521 4. 756 8. 494 85. 036 7. 960 16. 178 26. 975	23. 428 24. 695 36. 632 442. 120 3. 396 10. 637 52. 450 18. 122	19.757 10.791 32.279 272.494 8.814 10.973 34.852 12.612 32.094 27.825	13.795 21.900 26.903 21.625 5.401 5.252 86.629 10.591

Table 28

	(UCUGA 1002ACO	40.219	26.894	55.357	16.296	14.511	28.531	22.648	15. 420
	HEMBA 1003968			139.030	117.010	33.973	54. 122	29. 356	338. 820
	HEMBA1003974	147. 157	439.547	13. 159	9. 187	5.748	6.820	6.962	10.367
5	HEMBA1003976	20.167	17.809		6. 558	6.776	12.413	17.016	9.367
•	HEMBA1003977	32.761	12.350	24. 212		11, 203	23.881	20. 489	17.488
	HEM8A1003978	40.564	13.858	10.812	11.585	31.348	48.051	31, 355	42.728
	HEMBA1003981	65. 803	34. 462	71.399	26.801		3. 781	3. 102	64. 356
	HEMBA1003982	15. 104	89.360	20.946	18.086	1.620		5. 977	7.569
•	HEMBA1003985	15, 199	10.866	21.715	9.199	1.517	8.041		
40	HEMBA 1003987	48.695	30.080	108.473	25.632	23.222	28.008	21.302	24, 940
10	HEMBA1003989	47.841	51.466	128.889	32.288	24, 298	24.627	15. 392	23.174
	HEMBA1004000	36.424	35.098	34.843	16.292	19.541	20.604	16.803	21.872
	HEMBA1004006	8.411	42.393	12. 931	2.863	3. 395	0.000	4. 943	9.742
	HEMBA1004007	135. 300	114.014	286.000	90.971	64. 473	74. 153	71. 985	79.319
	HEMBA1004010	58. 331	152.845	38.785	18.676	18.819	35. 229	31.514	80. 599
	HEMBA1004011	62.306	16. 294	38. 336	12.356	13.756	29.883	26.091	7.986
15	HEMBA1004012	47.010	38.053	139.110	42.415	22, 159	34.340	27. 215	32.550
	HEMBA1004015	24.416	26. 249	27.372	12.243	13.962	25. 082	25. 133	12. 269
	HEMBA1004024	149.457	114.788	479.037	80.679	77.896	75.066	57. 366	93. 859
	HEMBA1004029	81.485	31.944	43.520	19.897	20, 191	38.768	36. 482	19.376
	HEMBA1004038	26. 629	15.823	19.708	12.109	7.832	14, 400	12.855	17.771
	HEMBA1004042	8. 177	10.678	12.830	6.612	11.484	7. 963	11.320	10.405
20	HEMBA1004045	24.675	30.855	37.128	20.069	23.538	15.509	17.299	17.447
	HEMBA1004048	95. 795	48, 977	78.760	36.608	40, 779	45. 132	47. 334	63.844
	HEMBA1004049	55. 947	543.954	47.428	49.034	19. 297	56. 209	23. 320	68.865
	HEMBA1004051	69.776	31.608	51.948	13.046	25.684	38. 632	30. 423	32. 553
	HEMBA1004053	29. 222	70.670	84. 481	24.394	15.007	23. 414	13.218	23, 973
	HEMBA1004055	39.564	23. 202	34. 928	8.151	5.353	28.619	15. 237	14. 807
25	HEMBA1004056	136, 121	122.072	413.353	75. 363	81.883	66. 439	41.004	85.794
	HEMBA1004060	17.642	11.826	29.995	9.507	4, 910	13.895	8.679	8. 388
	HEMBA1004051	17. 144	13.460	20.009	16.913	8. 228	14, 145	12.424	5.810
	HEMBA1004067	165.029	79.589	104.390	62.419	50.783	89.115	94.004	91.850
	HEMBA1004071	28.405	34.722	37.707	19.775	14.692	17.342	23.864	27.554
	HEMBA 1004074	128.445	51.388	148. 050	35.606	37.851	50.216	53.461	46. 373
30	HEMBA1004078	26. 126	14.714	20.940	9.721	16.211	17. 198	17. 388	14.057
50	HEMBA1004085	42.006	24.067	36.862	15.417	17.609	19.555	28.362	21.993
	HEMBA1004086	27.330	49.843	21.238	43.213	24. 232	16.260	12.409	22.262
	HEMBA1004097	45.296	15. 292	27.795	13.971	26. 928	26.002	33. 192	19.361
	HEMBA1004100	40.930	37.210	48. 942	23. 245	10. 184	25.744	21.452	28.594
	HEMBA1004103	101.036	101.281	184.668	64.176	44. 322	55. 385	41.050	40.000
	HEMBA1004110	89.903	65. 107	57.751	43.841	27.836	21.315	27.631	34, 280
35	HEMBA1004111	171.907	134.108	296.310	95. 474	115.874	78.450	80.011	98.760
	HEMBA1004124	177.408	71.838	103.065	37.865	46.198	68.531	109. 364	77. 083
	HEMBA1004130	64.543	54. 797	171.602		35. 382	25.601	19.599	23. 097
	HEMBA1004131	41.654	24. 184	33.975		23. 365	28.790	20.022	24. 999
	HEMBA1004132	55.906	42.840	162. 243		30. 251	28.863	19.780	22.237
	HEMBA1004133	64.624	30.838	38. 522		20.897	28.027	28.747	33. 333
40	HEMBA1004138	61.197	21.853			9.337	30, 080	17. 345	22.082
	HEMBA1004143	15.715	9. 656			10.539	14.067	11.441	9. 994
	HEMBA1004146	40.893	21.789			32.870	23.542	14.368	20. 982
	HEMBA1004148	59.990	18.796			17.531	18. 309	29.374	22.628
	HEMBA1004149	16.284	11. [3]	18.385			7.577	5. 890	13.683
	HEMBA1004150	5. 223				2, 553	2.158	2.062	2. 260
45	HEMBA1004154	111.110	40.836				58. 472	62.983	47.866
	HENBA1004164	139.670	107.565	315. 189	77.326		57.372	46.726	67. 257
	HEMBA1004168	24.042						3. 054	
	HEMBA1004199	22.894	9.047	10.46	8.631				
	HEMBA1004200	33.301		83.46				17. 235	
	HEMBA1004201	54.766		32. 37	0 17.449				
50	HEMBA1004202	14. 526			4 6.804				
	HEMBA1004203	47.655			2 13. 604			16.079	
	HEMBA1004207	6.344							
	HEMBA1004210					41.396			
	HEMBA 1004225					43. 156			
	HEMBA1004227						28. 177	25. 468	30.978
55	11000011004551								
JJ									

Table 29

			C7 144	62.536	27.672	34.345	69.613	47.182	38.807
	HEMBA1004235	99.954	57. 144		18. 289	11.697	19.212		16. 922
	HEMBA1004237	27.504	21.542		34. 130	27.841	36.089		34. 578
5	HEMBA1004238	79.210				1.072	2, 912	4, 422	1.294
3	HEMBA1004241	5.663	2.654	7. 035	2.556		85, 478		62.567
	HEMBA1004242	256.862			80.010	76.455			47.743
	HEMBA1004243	72.699	55. 276	60.764	28. 287	47.148	36.800		18. 700
	HEMBA1004246	44,915	30.967	100. 300	22.414	17. 109	15. 470		
	HEMBA1004247	66.750	16. 238	24.674	18.889	22.763	31.897	38. 415	17. 377
	HEMBA1004248	13.953	18. 412	17.581	11.953	11.378	14, 538	12.794	9. 562
10	HEMBA1004250	24. 439	10.494	10,631	6.401	5. 142	14.218	12.652	11.966
		37.349	20.650	22. 246	9.949	9. 550	14.570	21.841	18. 200
	HEMBA1004252	10. 994	19. 320	16.415	15.707	20.374	13.845	11.265	19. 838
	HEMBA1004260	10. 334	14.715	13. 358	7.615	5. 234	12. 282	15.089	11.397
	HEMBA1004264	22.716	195.750		171.071	174. 292	115.073	102.973	144. 125
	HEMBA1004267	235. 310		23.678	13.063	12.012	15, 529	14. 123	14. 593
15	HEMBA1004272	28.776	19. 025	53. 598	30.356	35.472	42.005	58.020	51, 617
13	HEMBA1004274	62. 157	50. 491		17. 443	18. 132	34.031	36.295	22. 171
	HEMBA1004275	70.423	38.514	45. 176	9. 548	9. 099	14, 035	10.406	8.615
	HEMBA1004276	33.630	4. 481	14.011		7. 333	10. 255	8.919	12.068
	HEMBA1004279	16.536	11.082	13.356	14.834	17.646	17. 327	17.630	13.770
	HEM8A1004284	29.688	30. 297	64.483	13.658		17.625	23.811	13. 547
	HEMBA1004286	32. 471	16. 566	18.049	12.391	6.773	37.379	28. 939	41.740
20	HEMBA1004289	81.573	62.930	165. 571	49.704	34. 785	57.764	53.695	45.065 -
	HEMBA1004293	72.465	34. 902	48.669	32.705	17. 408		23.797	20. 926
	HEMBA1004295	37. 595	12.116	29.975	11.634	5.514	25.018		9. 355
	HEMBA1004302	10.880	5.912	7.885	10.025	5, 190	6.060	5. 264	
	HEMBA1004306	426.811	177. 321	335. 168	107.646	123.947	256.397	251.772	134. 005
	HEMBA1004312	37. 953	30.864	105. 533	30.747	25.847	16.140	16. 283	24. 272
25	HEMBA1004314	29. 396	23.332	95. 584	22. 179	18.544	11.015	8.804	20. 974
23	HEMBA1004321	47.670	29, 150	105, 316	35.655	23. 139	31.309	29.736	47.858
	HEMBA1004323	87. 295	65. 931	221.440	44.690	41.425	36.609	34.117	39. 135
	HENBA1004327	65.869	21. 284	21.540	11.985	14.419	27.213	27.030	20. 118
	HEMBA1004329	67.920	44. 687	132.755	32.977	21.556	32.356	17.478	26.773
	HEMBA1004330	8.765	7.655	16.827	7.164	3.843	9.511	7.660	4.615
	HENBA1004334	16. 438	21.355	31.580	15. 109	26.670	13.368	10.581	13.568
30	HENBA1004335	204. 961	102, 859	325. 226	69.979	64.392	78.772	71.641	83. 525
		186.677	30. 208	51.439	15. 995	41.404	87.221	89.558	40. 224
	HEMBA1004341		76, 316	123. 332	42.705	51, 432	26.797	42.054	59.071
	HEMBA1004344	261.676	32.610	97.858	37.038	21.953	33.115	33.526	36.846
	HEMBA1004347	65.249	35.727	29. 441	19.803	18.786	23. 126	19.103	18. 719
	HENBA1004349	22.353	65. 544	237.050	49. 039	34, 141	32.597	28.166	46.343
35	HEM8A1004352	75. 508	66. 042	132.169	40.563	27.380	39.551	30.556	56.886
	HEMBA 1004353	54. 322	29. 352	79. 264	22.784	20,533	21.755	16.860	22.429
	HEMBA1004354	43.687	22. 201	27.487	10.404	8.280	22. 159	16.039	15.038
	HEMBA1004356	44.730	28, 429	71.634	26. 232	36, 259	59.602	38, 361	50.410
	HENBA1004360	91.412	10.099		5. 481	5, 631	6.802	6,791	6.167
	HENBA 1004365	9.956		5. 338		1.638	1.507	3. 555	1.568
40	HEMBA1004372	3.613	4. 593			26.163	34. 545	30.827	33. 572
40	HENBA1004377	53.834	41.410			13, 968	16, 520	14. 951	17.114
	HEMBA1004389	20.540	22.800			19.018	31.716	23, 931	23.617
	HEMBA1004391	60.284	22.653			75. 399	76.587	39.777	44.665
	HENBA1004393	177.786	197. 548				17. 825	10.981	8. 836
	HEMBA1004394	28. 949	11.849			16, 519			21.980
	HEMBA 1004396	17.907	26.956					20.530	20.774
45	HENBA1004401	22.519	21.858		_				29.842
	HEMBA1004405		38. 835				19. 522	18 846	20.863
	HEMBA1004408	50.497	27.15						36. 385
	HEMBA1004414								46.317
	HENBA1004429	61.867							42.005
	HENBA1004433		39. 82		37.521				35.073
50	HENBA1004440								37.635
	HEMBA1004444		46.58						
	HEMBA1004446			9 29. 42					16.078
	HEMBA100445								
	HEMBA 100445			8 18.47	9 3.44				
	HEMBA1004454				6 24.38	2 20.63	24. 49	4 22.897	29.042
66	III.ADA 100443								
5 <b>5</b>									

Table 30

				1 20	Die 30				
	HEMBA1004460	138, 550	96. 143	356.058	74.883	54.735	70.698	38.344	55.945
	HEMBA1004461	64.074	19.163	24.808	12.846	16.373	23. 508	22.827	10.137
	HEMBA1004468	134.439		210.409	77.409	60.142	56.229	42. 361	49.457
	HEMBA1004479	82, 994	32.899	92.282	35. 519	44. 435	43.183	31.836	37.116
	HENBA1004482	5, 602	7.682	11. 248	36.034	2.926	5. 535	5.693	5, 972
	HEMBA1004491	16. 736	6, 285	17.615	14.018	11.729	19.804	10.683	11.280
	HEMBA1004499	94, 095	71.456	148. 355	58.479	48.596	46, 968	46.648	57.279
	HEMBA1004502	21.523	9.344	18. 265	7.282	9.979	2.762	3.174	13.389
	HEMBA1004505	26,042	15. 980	43. 855	17.516	15. 469	22.190	22.873	15.812
)	HEMBA1004506	12.004	29. 395	42.664	21.849	21.426	14. 459	9. 224	9.845
	HEMBA1004507	96, 377	87.688	99. 177	103.472	34, 160	81.068	54.939	151.142
	HEMBA1004509	52.657	14.880	19. 120	16. 228	17.009	24.783	24.565	13.476
	HEMBA1004523	20. 156	18, 209	11.197	16.529	14.651	13.004	20. 267	19.467
	HEMBA1004528	42,620	27.819	48.069	14. 426	25. 267	43.038	40.239	37.718
	HEMBA1004514	75.090	41, 159	44. 399	31.300	16.686	31. 317	21.009	18.589
5	HEMBA1004536	31.531	13.343	23.664	14.085	4. 408	13. 133	12.981	15. 201
-	HEMBA1004538	352.363	181.508	233.819	97.018	122.402	183.507	100.197	150, 062
	HEMBA1004542	47, 360	17, 733	29. 238	17.280	12.324	17. 317	22.764	15. 212
	HENBA1004552	63.401	29. 585,	26.857	43.567	39.674	38.686	23.830	33. 542
	HEMBA1004554	62.231	11.953	25.084	16.273	32.820	33.809	20.812	5. 536
	HEMBA1004558	30.217	12.133	31.036	15.840	30.638	65. 183	19.155	30. 921
)	HEMBA1004560	68, 901	16.566	17.908	16.431	10.034	30.093	22. 379	22.683
,	HEMBA1004564	48.119	14.911	35.565	31.983	32.464	30.028	20.965	32.479
	HEWBA1004566	32.479	29. 553	20.970	32.788	42.949	40.715	23. 273	32. 960
	HEMBA1004573	17. 728	13.843	7.118	9.972	19.952	9.755	9. 278	8. 100
	HEMBA1004576	39, 572	26.733	42.044	10.704	37.505	32.441	17. 232	25.36
	HENBA1004577	46.233	11.570	97.881	39.434	13.437	41.089	34. 426	35. 314
-	HEMBA1004586	82.532	71.398	213.814	70.289	52.589	45.729	23. 395	38. 31
5	HEMBA1004596	72.534	32.493	45.820	27.585	27.854	34. 997	33.847	38. 47
	HEMBA1004604	99.019	48. 582	103. 587	36.723	49. 392	48. 377	56.558	69.25
	HENBA1004607	53, 557	37.013	100. 999		26.143	28.796	21.692	42.04
	HEMBA1004510	20.690	14.854	69. 908		12.120	9, 108	8.858	15.08
	HEMBA1004617	22,592	20. 386	42.426		15, 568	10.691	6.697	10.31
_	HEMBA1004622	78.025	46.803	209.059		29, 836	29.902	12. 194	27.43
10	HEMBA1004626	38.170	36. 312	110.684		14.118	17. 193	15. 579	20.82
	HEMBA1004629	33.858	37.886	87.440		47.341	28. 160	9. 569	32.85
	HEMBA1004631	35.946	10.475			17.128	22.775	22. 566	6.36
	HEMBA1004632	27.084	13.891	23.598		7.802	11.754	48. 857	40.81
	HEMBA1004633	78. 391	33. 135	114.054		49.008	60.659	13. 502	25. 03
	HEMBA1004636	52.397	20.706	34.962		22.609	21.255	2. 597	5.02
35	HEMBA1004637	4. 228	4.304			9.756	4.086	0, 113	0.00
	HEMBA1004638	0.241	0.000				27.214	20. 560	24.84
	HEMBA1004645	57. 971	29. 263				14.099	17, 122	12.00
	HEMBA1004656	16. 139					42.697	9.048	13.3
	HEMBA1004657	20. 820		_			5. 098	2.525	7.5
	HEMBA 1004666	7. 321	3. 174				28.018		25.6
10	HEMBA1004669	94.910						23.413	14.3
	HEMBA1004670	57. 231						20.328	
	HEMBA1004672	63.471							
	HEMBA1004689								
	HEMBA1004690								
	HEMBA1004693								
45	HEMBA1004697							4 4 44	4 47 6
	HEMBA 1004702								
	HEMBA1004704								
	HEMBA1004705								
	HEMBA1004706								
	HEMBA1004709								
50	HEMBA100471								
	HEMBA100472:								
	HEMBA100472								
	HEMBA1004730								
	HEMBA100473:	30.76	9 29.88	23.3					
	HEMBA100473		2 11.97	4 36.5	95 3.98	18 12.55	6 7.65	3   4.30	3 1 12.

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Table 31

		C	***	T-2 -12 1	48 681 T	38.696	23.516 1	19.970	34, 509
	HEMBA1004736	55. 309		132.333	45.653			12.665	18. 349
	HEMBA1004748	53.832		172.069	29. 562	22. 161	14. 904		42.250
5	HEMBA1004749	127. 285	45. 137	73.698	27.788	33. 184	60.214	44.635	
•	HEMBA1004751	81.283	64. 830	173.888	54, 165	36.368	41.802	31.283	43.505
	HEMBA1004752	59.058	32.785	109.428	32. 254	29.090	34. 259	30.970	33.029
	HEMBA1004753	204.044	247.465	406.165	156.689	102.755	131. 323		263.882
	HEMBA1004755	57.638	59.677	83.850	22.148	29.800	30. 642	13.064	23. 261
	HEMBA1004756	9. 965	16. 228	11.023	8, 349	6.780	9. 109	111.628	14.885
	HEMBA1004758	36.487	26. 558	116.970	22, 341	14. 553	14.773	11.840	14.406
10		67. 343	19. 641	33.742	13.841	16.720	25. 489	23.061	18.650
	HEMBA1004763	29. 177	24. 043	38.303	6.673	10.298	3. 197	10.352	13. 391
	HEMBA1004768	10. 327	14. 492	10. 901	6.416	6.310	7.963	10.868	7.955
	HEMBA1004770		34, 314	76.491	31,609	22.830	23. 102	30, 433	32.358
	HEMBA1004771	46.910	28. 706	63. 968	24, 931	18.754	43.049	32,720	26.795
	HEMBA1004775	39. 253	11.017	10. 103	5. 466	9.000	16.400	10.105	8.046
15	HEMBA1004776	22.604		223. 475	37, 540	33.791	32. 337	24.067	43.529
	HEMBA1004778	78.144	77. 681	102.812	8.313	15. 151	11. 373	9.479	6.329
	HEMBA1004784	9.826	18. 370		6.651	10. 549	10.674	13.732	11.615
	HEMBA1004785	25. 723	16. 345	26.216 16.096	7.804	8.691	10.011	7,713	11.389
	HEMBA1004789	18. 173	14. 5081		11.028	9.351	9.757	9, 905	12.028
	HEMBA1004795	14. 283	12.973	25. 122	34.142	28. 246	40.067	32.715	25, 583
20	HEMBA1004797	65. 927	33.745	73.888	27.072	30. 957	22.607	22.520	26.554
20	HEM8A1004803	36.634	41.124	65.880	8.868	9.653	9. 000	7.894	8. 199
	HEMBA1004806	11. 997	8. 183	21.467		12.009	13.340	7.935	9.118
	HEMBA1004807	16. 352	14. 481	22. 459	11. 249	29. 259	8. 180	12.578	10, 934
	HEMBA1004816	29. 782	24.075	95.884	18.110	4.018	4. 269	6.876	3.493
	HEMBA1004820	8.636	7_466	8. 862	4. 249	64.754	83. 155	56.657	65, 121
	HEMBA1004833	159, 947	50.729	81.248	38.650		35.053	31.537	40.529
25	HEMBA1004847	51. 456	25. 570	40.694	21.115	36. 572 26. 080	54.413	50. 197	24. 185
	HEMBA1004850	77.254	24.014	38.620	21.854	31.887	25. 058	20.050	20.982
	HEMBA1004863	57.117	32.704	72.480	23.951	13.898	16.719	20.308	17.843
	HEMBA1004864	46.043	27.344	59.824	26.750	14. 331	13. 454	13.835	15.797
	HEMBA1004865	12. 257	14. 642	31.748	44.090		24.064	20.647	23, 264
	HEMBA1004880	56. 788	50. 021	126.837	35. 420	26. 589	13.652	10,676	19.977
30	HEMBA1004882	42, 450	18. 453	29.340	16.782	13.013	3.711	3, 652	6.615
	HEMBA1004885	8. 545	4.947	5, 350	4. 891	2.933	23. 391	15.605	16.916
	HEMBA1004889	28. 103	22.485	32.049	17.078	14. 363	6. 539	5. 245	9, 440
	HEMBA1004900	19. 922	15.709	33. 254	10. 423	9.045	36.045	18.861	27.933
	HEMBA1004909	88. 522	49.269	163. 284	48. 147	35. 537		15.857	25. 370
	HEMBA1004918	64. 384	43.134	105.868	34.899	22. 323	18, 192	10.213	20.111
35	HEMBA1004923	47.731	37.996	69.168	19.659	26. 441	9. 882	8. 967	11.809
00	HEMBA1004929	11.048	14.003	10.808		7.539	34. 331	31.091	41.874
	HEMBA1004930	101.277	92.425	279.652	80.664	12.296	10.327	96. 467	5. 417
	HEMBA1004933	9.145	5. 566	12.895			5.866	9, 143	12.805
	HEMBA1004934	7.311	7, 106	43.966		4.750 17.734	15. 280	15.784	46. 365
	HEMBA1004937	43. 331	27. 219	38.802			27.585	29.628	38. 533
40	HEMBA1004943	51.072	26. 833	32.001			38. 514	31.589	23.074
40	HEMBA1004944	84. 363	46. 788	126.294			27.854	34.636	31,712
	HEMBA1004946	64. 538	28. 144	37.908			33. 568	20.062	19.020
	HEMBA1004952	90. 835	18. 893	40.862			20. 777	6. 946	29. 261
	HEMBA1004954	14.656	36.003	41.485			4. 578	1. 565	5.188
	HEMBA1004956	5. 975	9. 923				47.560	29.646	45. 929
	HEMBA1004960	86.030	77. 420					19.705	18, 480
45	HEMBA1004971	31.046	5. 439					30.004	50.710
	HEMBA1004972	77.318							22.580
	HEMBA1004973	35. 524							10.523
	HEMBA1004977	6.756							10, 106
	HEMBA1004978	8.689							22.124
	HEMBA1004980	34.093							5. 456
50	HEMBA1004982	14.750							8.305
	HEMBA1004983	38. 285							33.683
	HEMBA1004995								14.698
	HEMBA1005004	13.855							16. 940
	HEMBA1005008								
	HEMBA1005009	34.543	15. 67	19.46	4 10.04	17.12	20. 30		

Table 32

	DIFTIO LANCOTA I	40 350 1	24 072 (	25, 349	20.834	30, 144	32.629	20.777	27.016
	HEMBA1005019	49. 260 37. 224	24. 872	38, 065	17.061	15. 647	20.369	25.859	32. 656
	HEMBA 1005021		63.713	35. 352	16. 531	19.588	26.517	15.798	16.604
5	HEMBA1005029	30. 265	200, 167	574. 746	133. 872	210.689	177.872	156.563	170.510
	HEMBA1005035	393. 404	41.961	73.015	39. 541	44. 451	66.623	55.833	51.349
	HEMBA1005036	115. 345	19.922	57.018	13.971	11. 999	26. 427	15. 206	7.350
	HEMBA1005039	28.850 93.995	31.868	\$4.335	18.576	28. 338	31.562	31.930	23. 751
	HEMBA1005047	78.015	41.690	73.330	29.830	26. 504	35. 887	21.640	35.653
	HEMBA1005050	23. 050	15.803	29. 553	15.707	7.836	15.618	19. 435	13. 336
10	HEMBA1005062 HEMBA1005066	10. 980	11.364	31.553	13.509	5.668	10.541	5.005	10.849
	HEMBA1005067	39. 308	34. 578	39.795	44.519	24. 643	21.272	19.379	20. 121
	HEMBA1005070	73. 155	34, 949	68.556	29.956	38.004	38.211	48.007	31, 733
	HEMBA1005075	88. 089	37.798	148.675	40, 537	33. 271	33.074	28.661	30.201
	HEMBA1005078	100.064	37.746	66, 827	33.115	41, 170	55. 560	51. 231	17. 112
	HEMBA1005079	137.757	86, 238	294, 118	73.304	76.035	75.084	47.255	76. 170
15	HEMBA 1005083	18. 102	7.642	17.087	6.711	6.184	8.675	9, 287	9.609
	HEMBA1005084	82.712	38.248	47.063	26.664	27.435	37. 552	38.419	28.349
	HEMBA1005088	31.610	22.435	76.774	22.700	18.926	23.875	8.895	12.447
	HEMBA1005089	68. 944	55.156,	178, 226	34.742	32.350	38.645	22.869	28. 148
	HEMBA1005090	148.861	86.156	117.997	94.811	57.034	81.098	54. 187	116.066
	HEMBA 1005096	83. 125	30.911	63.940	33.378	33.962	48. 589	35.467	36.021
20	HEMBA1005101	69.080	14.020	34. 136	10. 197	13.998	34. 420	22.696	15. 975
	HEMBA1005107	82.659	25. 203	36.223	11.215	21.514	32.720	25.972	21.337
	HEMBA1005113	7. 977	17. 225	31, 501	7.563	44.493	5, 157	6.957	9.761
	HEMBA1005123	173.637	77.260	555. 672	126. 908	94.628	90. 446	70.735	90.016
	HEMBA1005133	58. 192	40.749	122.920	29.864	16.700	17.652	8.802	18. 988
	HEMBA1005135	8. 259	9, 125	14.962	2.213	16.732	6.892	3. 383	6.189
25	HEMBA1005145	185. 299	101.220	352. 159	92.082	88.750 125.192	122.118 96.704	76.475 92.083	90. 044 128. 030
	HEMBA1005149	220. 122	109.352	274.492 226.882	120.663 58.505	33.738	46.323	27. 534	34. 457
	HEMBA1005152 HEMBA1005159	125. 948 15. 760	96.291	9. 399	6. 198	6. 191	6.861	12.001	4, 556
	HEMBA1005172	1653. 208	89.658	73.666	54.667	33.118	55.680	32.520	70. 907
	HEMBA1005185	9. 954	17.248	10.492	34. 452	3. 558	3, 117	6.026	11.173
••	HEMBA1005186	23.745	10.048	27.091	13.067	7.719	15.412	15.086	15. 591
30	HEMBA1005195	14. 573	8.648	11.038	19.306	6.313	25.313	13.510	9. 183
	HEMBA1005201	52. 322	13.197	47.505	13.091	12.078	8. 531	23. 532	9.848
	HEMBA1005202	98. 566	30.141	71.588	27.954	44. 381	66.294	42. 390	39.695
	HENBA1005204	184. 429	287.156	382.039	168.753	203. 458	222. 970	143.609	358.646
	HENBA1005206	148, 946	61.309	84.791	34, 139	49.115	66. 295	83.608	76.159
35	HEMBA1005219	21.685	17.755	9, 606	8. 236	8.038	13.031	7.751	11.441
33	HEMBA1005223	79.969	42.665	113. 460	40. 547	32.099	53.017	26.025	32.004
	HEMBA1005229	26.819	9, 926	21.841	3.135	5.090	6.656	4.681	7.079 65.511
	HEMBA1005230	71.184	67.313	201.065	79. 279 8. 552	59.679 3.285	77. 484 12. 098	4.975	3.965
	HEMBA1005232 HEMBA1005238	7. 374 96. 780	6.386	17.522 51.932	8. 128	20.776	69. 291	49.474	35.019
	HEMBA1005241	142. 598	104. 185	428.635	78.773	78.033	74.434	42.333	63.097
40	HEMBA1005244	76.771	32.597	37.797	16.459	12. 489	35. 934	31.814	35.602
•	HENBA1005246	241. 316	60, 348	73.077	25.067	41.351	117.666	88.193	54.014
	HEMBA1005251	37. 505	33.247	108. 631	23.585	14, 915	23. 393	14.302	16.409
	HEMBA1005252	53. 401	25.532	37.199	15.002	20.744	31.279	24.207	27.562
	HEMBA1005267	17. 238	39.564	20.097	27. 506	31.874	11.013	14. 526	14. 024
	HEMBA1005274	16. 538	8.744	18.308	9.021	10.103	11.943	8.914	11.978
45	HEMBA1005275	69.133	43.329	215.468	46. 290	57.647	37.411	25.040	41.913
	HEMBA1005288	65. 401	50, 495	150.714	33.833	34.633	28. 241	24. 910	40.164
	HEMBA1005293	17.403	9, 430	23. 201	4.467	3. 192	25. 520	6.775	8.771
	HENBA1005296	223. 097	811.623	894.835		220. 523	698.319	418.435	1376. 785
	HEMBA1005301	36.708	16.970	29. 798	11. 929	12.544	22. 221	35.726	32.270
	HENBA1005304	83.978		260.016	50.686	36.101	36. 160	24.896	47.838
50	HENBA1005305	44. 218		74.215	27. 494	27.352	34. 920 8. 958	21.424	38.882
	HEMBA1005311	33.034		48.263		7.908	14. 209	9.090 63.072	9. 124
	HEMBA1005313	11.165 6.948	36.175	17.550		3,812	5. 787	4.069	4.891
	HEMBA1005314 HEMBA1005315	72.349		156.842	34.545	43, 132	26.415	28. 942	28, 442
	HEMBA1005317	20. 230		17. 258	9.796	11.664	9. 976	9. 263	8.017
	INC. MONTO 03317	20.230	0. 431	17. 230	3.,50	1	1		
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Table 33

	CUENTA LANGESTA		A44 T	12 663 [	5. 228	5.376	9.013	5, 511	4.846
	HEMBA1005318	14.755	5. 931	13.883					17.915
	HEMBA1005324	98.070	33.348	44. 270	26.052	35. 446	48. 523	30.889	
_	HENBA1005331	24.826	335. 211	15. 947	26. 496	14.744	21.427	16. 942	29. 580
5	HEMBA1005337	19,080	18.022	19, 429	5. 217	20.830	29. 867	32. 481	44. 585
	HEMBA1005338	61.533	38.788	63, 113	23.657	30. 437	48. 455	40. 921	36.285
	HEMBA1005344	384.481	88. 937	143, 574	53.983	72.524	167.620	135. 992	68.042
			68. 949	220. 401	62,090	53.484	67.048	30. 456	42.612
	HEMBA1005353	111.629			59.707	36.743	34. 233	21.666	47.596
	HEMBA1005359	87.635	64. 332	175. 543			19. 092	20. 883	12.720
10	HEMBA1005362	25.674	25. 093	18.642	30.797	21.917			
10	HENBA1005364	5.677	2.817	5, 168	13.116	19.753	5. 180	2.877	7. 198
	HEMBA1005367	51. 911	28. 536	74.559	28.446	30.138	27. 987	16.766	22.415
	HEMBA1005372	11.289	5.819	11,700	5,659	9. 177	6.402	9. 312	4. 913
	HEMBA1005374	64.639	57.505	120, 218	32.738	30. 987	24.792	23.695	30.728
	HEMBA1005379	29.549	13.813	12.040	8.862	7.648	11.978	9. 051	4.019
	HENBA1005382	140.116	94.743	104.609	70.213	26.226	53.452	88. 235	85. 480
15	HEMBA1005384	33. 109	15, 221	21.713	10.250	8.543	11.030	7. 498	9.010
				52. 790	29.541	31.691	44.619	35, 179	29, 136
	HEMBA1005386	111.062	30.547			35.894	16.765	11.513	24.601
	HEMBA1005389	66. 821	32.429	129. 272	42.528			16. 944	24. 443
	HEMBA1005394	35. 794	18. 327,	22.715	25. 833	26.639	30.857		
	HEMBA1005403	40. 404	14.030	54.041	14. 621	15. 504	27.461	15. 586	32.390
	HEMBA1005408	51.701	45.069	71.813	44. 257	67.383	35.010	23. 690	44.612
20	HEMBA1005410	4.534	4. 269	11.774	12.035	10. 197	6.188	3. 955	8. 910
	HEMBA1005411	75. 220	94.039	163.001	67.133	50.499	41.243	22.652	35. 008
	HEMBA1005423	35.745	26.430	69, 138	35.773	15. 442	19.286	14. 057	23.010
	HEMBA1005426	14.366	12.073	14, 418	5. 345	11.591	8.954	3.082	7. 203
	HEMBA1005427	66.444	99, 596	61.088	47.865	59. 821	53.861	25. 223	46.397
	HEMBA1005430	52.945	15. 385	36, 316	19, 210	23.854	37.895	19. 556	18.127
	HEMBA1005438	51.806	28. 359	33, 314	17.787	19. 295	21.754	13.422	29. 941 .
25	HEMBA1005443		165.667	426.408	91, 550	77.559	76.024	105.042	108. 232
		108. 954	39.578	65. 244	29.171	28.000	21.457	18.763	23.755
	HEMBA1005447	51. 383		41.861		27.647	39, 311	28. 567	27. 508
	HEMBA1005449	86.452	20. 253		15, 939		53. 326	67. 529	72. 233
	HEMBA1005452	110. 567	52.128	74,119	42.532	39.847			11. 445
	HENBA1005454	7.997	16.821	17.998	14. 293	14. 436	8.454	6. 498	
30	HEMBA1005468	185.066	78.008	126.372	56,026	56.490	78. 922	61.083	57.511
30	HEMBA1005469	88. 419	54.761	196, 280	63.682	53.661	42.639	23. 441	30. 144
	HEMBA1005472	37.878	41.710	88. 807	34, 196	28. 126	21, 983	24. 350	30. 575
	HEMBA1005474	89.169	55. 263	212.086	51.664	50.480	66.508	39. 590	30. 322
	HEMBA1005475	212.273	98.359	182.707	110.945	105.968	98.316	56. 095	68.647
	HEMBA1005489	61.603	40, 439	42.459	21.361	21.335	31.130	11.578	25.898
	HENBA1005497	10. 325	12. 398	5, 705	8. 252	5.611	9. 949	2.213	16.039
35	HEMBA1005500	86.636	39, 755	180.843	46.031	28.664	31.809	14.951	31.189
	HEMBA 1005506	24. 029	3. 468	17.794	7.400	5. 464	9.419	6. 593	3.517
	HEMBA1005508	12. 944	12. 524	22. 247	8.535	16.857	11.561	7,741	12.059
	HEMBA1005511	116. 338	59. 193	267.636	59.921	58. 995	43, 190	30.786	51.049
			70.217	88.519	56.620	54. 920	73.797	80.751	68. 624
	HENBAT005513	167. 332		23.901	8. 903	17.777	21.966	23.844	18.611
	HENBA1005517	37.667	10.443			36. 856	47.397	27. 618	27.825
40	HENBA1005518	109. 105	25.679	71, 345	23.319	106.207	95,070	67. 199	94.086
	HENBA1005520	200. 267	104, 176	459. 373	133. 255			16.899	
	HEMBA1005522	36. 421	15, 946	24.796	12.598	8.472	14.558		13.857
	HEMBA1005526	116. 274	72.899	292. 397	82.002	73.603	66.198	34. 319	47.682
	HENBA1005528	13.037	9.406	30.550	14.612	15. 947	16.516	7, 583	24. 988
	HEMBA1005530	56.516	26, 583	63.811	13.686	21.441	29.159	24. 254	21.717
45	HEMBA 1005538	5. 523	17.373	36. 952	7.017	10.885	11.406	15. 411	35.789
45	HEMBA1005539	76.498	30.847	69, 424	17.584	24.989	35. 829	28.772	25.913
	HEMBA1005545	46. 912	10.940		12 22	46.822	33.595	31.865	24.090
	HENBA1005548	57.779	14. 326	15.050		18.538	22, 115	22. 271	39. 291
			120.695			81.893	79. 223	60.281	62.088
	HENBA 1005552	141. 489		24. 397		22.919	24.422	21.466	8. 178
	HEMBA1005558	52. 488				38.261	33.077	24.038	37.014
50	HEMBA1005568	74. 152	61.206				31. 212	27. 080	30. 221
	HEMBA1005570	54. 151	68. 747						19. 991
	HEMBA1005575	71.454				8. 931	30. 461	29. 371	
	HEMBA1005577	40.771	13.448			6.610	18. 266	12.838	10. 181
	HEMBA1005581	81.577	27.270	38, 708			33. 479		16.842
	HEMBA1005582	24. 681	30, 135	30. 933	14. 220	7.764	10.454	12.847	13. 157

443

Table 34

	HEMBA1005583	23.564	22, 466	98.629	9.735	10.545	12.468	10. 523	17.884
	HEMBA1005588	96.041	96. 264	266.022	69.126	54. 588	44.105	34.310	52.441
				125.688	37.987	35. 953	41.577	39.834	47.357
5	HEMBA1005593	51.102	40. 350			8, 750		12.433	7. 264
	HÉMBA 1005595	52. 429	18.652	31.240	8.095		14. 586		
	HEMBA1005597	125.119	43.335	90.414	24, 402	44.780	74.946	66.352	45. 322
	HEMBA1005606	141, 646	65, 667	95.041	30,084	57. 974	107. 329	84.655	46. 145
	HENBA1005609	77. 991	60.190	244. 951	52.002	41.602	40. 406	26.928	42.614
					42.670	41.809	32. 256	23. 683	43. 139
	HEMBA1005616	47.760	62.865	190.870					
10	HEMBA1005621	33.797	18. 993	22. 515	11.333	11. 545	16. 964	12. 122	13. 910
10	HEMBA1005627	128. 661	66.487	148. 021	45.359	42.161	42.054	30.884	43. 319
	HEMBA1005628	43.539	36.758	85.714	25.524	46, 601	19. 229	82.784	36.636
	HEMBA1005631	21, 340	8. 467	38.068	22.476	18.318	17.813	12.599	28. 199
				233.637	59.097	45. 388	52.090	29. 944	37. 461
	HEMBA1005632	113, 190	73.651			107. 528	72, 729	54. 939	130. 473
	HEMBA 10056 34	123,668	195, 912	390.579	101.523				
	HEMBA1005662	15. 391	11.345	23.021	7.453	5. 561	13.084	8. 973	5. 282
15	HEMBA 1 005666	33.844	30, 419	34.983	13.220	31.573	24. 609	13.796	28. 043
	HEMBA1005670	91.667	63, 609	255. 523	57.730	46.927	45. 285	23.794	46.684
			55. 388	34.948	26.297	20.567	2.367	5.666	13.509
	HEMBA1005671	63, 448				32. 151	40.446	37. 522	36.817
	HEMBA 1005679	53. 089	33. 284	126.705	39.666				
	HEMBA1005680	115, 289	72.018	220. 408	76.653	55.707	68.735	32.613	36. 282
	HEMBA1005685	68, 783	45.211	72.197	30.110	32.724	43. 022	37.740	33.510
20	HENBA1005698	37.890	35.679	44.793	29.794	25. 150	48.613	15.651	20.648
	HEMBA1005699	14. 243	17.539	37. 269	9.035	12.276	5. 454	5. 259	6.787
				20. 249	7.662	15. 489	11.648	8.488	10. 229
	HEN8A1005703	19.524	15.116			29.605		18.303	44. 730
	HEMBA 1005705	35. 316	35. 677	66.552	26.492		90. 298		
	HEMBA1005712	20, 312	29.695	30. 267	17.829	17.668	17.695	14, 517	23.820
	HEMBA1005717	47.313	15.037	30.499	6.950	13. 391	32.044	15.084	7.078
0.5	HEMBA1005718	88.576	81.734	175.773	75.414	46.080	58.797	49.803	76.705
25	HEMBA1005721	84. 981	42.340	58. 434	18.134	34. 245	43.284	34.523	41.460
			92, 346		55.652	48.768	63. 471	92.755	56.031
	HENBA1005722	174, 952		194.868				9.324	5. 953
	HEMBA1005724	32.655	8. 284	5. 342	4.000	14.801	15. 671		
	HEMBA 1005732	89.624	24. 907	32.546	5.638	21.753	30.046	28. 487	20.595
	HEMBA1005737	25. 179	16.797	16.017	10.703	12.731	12. 444	8. 579	7.257
	HEMBA1005742	11.547	23.162	24. 345	20.921	29.934	18. 597	13.749	22.702
30	HEMBA1005746	36.098	14.407	21.907	16.923	13. 431	12.235	10, 908	8.606
					21.861	30.274	80, 588	47.082	26.037
	HEMBA1005747	80.718	30.396	44. 843					
	HEMBA 1005749	35.749	31.758	64.769	22.766	28.853	25.733	31.698	30.753
	HEMBA1005755	34.680	39. 133	30.663	37.837	21.308	24. 392	15. 905	25.470
	HEMBA1005760	118.125	41.490	33.276	25.724	28.933	46. 295	36. 173	31. 205
	HEMBA1005765	94, 451	70.516	200. 826	48.023	37.340	35.414	31.098	40.041
35	HEMBA1005766	112.861	70.359	87.247	48. 958	51.073	52.147	72.391	63. 859
					28.456	45. 254	28. 283	25. 156	29. 122
	HEMBA1005780	55. 961	34.713	89.816					
	HEMBA1005795	18.800	38. 386	19.666	10.007	13.009	11.811	13.106	14. 493
	HEMBA1005809	67.301	66.510	87.390	53.061	43. 975	35.574	35.334	57.818
	HEMBA1005813	52.911	84.881	160.064	38.752	43.727	30.799	23.426	57.177
	HEMBA1005815	30, 398	30.434	43.366	19.911	16. 123	39.746	26.743	28.548
40	HEMBA1005822	40.948	47.746	65. 298	51.932	30. 845	20.187	22.641	29.114
40				272.004	48.816	36. 558	40. 259	23.443	35. 824
	HEMBA1005829	114.982	70.536						18.807
	HEMBA1005833	59. 540	25.743	29. 266		24.711	26.964	17.968	
	HEMBAT005834	151.440	82.917	322.413		74.711	59. 590	35.082	70.415
	HEMBA 1005844	66.624	11.865	96, 556	95.719	56. 133	75, 546	55, 974	122.840
	HEMBA1005852	71.743	77.830			85.623	78.593	90, 291	87.310
	HEMBA1005853	62.809	83.326		63.897	79. 208	48, 939	27, 359	58.468
· 45						65. 325	53.917	35. 383	63.446
	HEMBA1005878	139.991	109. 928			1 2 2 2		100	11 000
	HEMBA1005883	5. 211	6.310			10.070	6.635	4.486	11.850
	HEMBA1005884	9. 136	10.768	29. 442	9.504		9.142	4.561	12. 287
	HEMBA1005891	8. 927	12.500		5. 996	7. 370	7.346	1.250	5.470
	HEMBA1005894	70.006	59.347			29. 584	23. 227	14.651	36.934
								23.019	41.130
50	HEMBA1005898	84. 199	61.254					26.778	43.681
	HEMBA1005902	38.306	16.873						
	HEMBA1005907	4.806	3. 997					5, 311	4. 941
	HEMBA1005909	4, 140	3.733	23.479	2, 443	4.661	6.683	0.750	
	HEMBA1005911	143.926	92.633			51.954	60.593	39.302	55. 189
	HEMBA1005912	18.801	17.269					12.917	
	INCHOMINGSTA	1 10.001	1 11.603	, , ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	, ,				

444

Table 35

								4 4 4 4	
	HEMBA1005913	10. 533	16.117	14.368	16.655	8.179	7.135	7.907	12.918
	HEMBA1005921	83, 262	45. 648	252.573	51.044	41.764	22. 286	23.762	46.202
					20.084	33.779	24.835	18. 394	14.883
5	HEMBA1005922	64.440	17. 427	35, 136					
•	HEMBA1005929	173.002	139.696	378.444	95.543	83.075	72.298	55. 205	94.716
	HEMBA1005931	146.354	89, 551	224.601	83.623	63.406	73. 122	54. 973	59.891
			91, 791	227.012	89.834	99.341	96.876	62.967	55. 492
	HEMBA1005934	141.558							
	HEMBA1005945	144.693	21.871	38.980	19.915	46.699	78, 590	80.430	30.052
	HEMBA1005962	67. 209	34.719	63.745	21.004	17.931	29. 331	21. 199	20.008
		18.320	6. 954	9. 127	5.913	2.497	8.674	7, 574	4.873
10 .	HEMBA1005963								
10	HEMBA1005990	581.646	117. 336	139, 967	53.671	242.262	424. 182	418, 873	85. 511
	HEMBA1005991	67.437	59. 327	188.570	42.994	21, 101	33.868	19. 164	_11.619
	HEMBA1005999	193, 878	135. 695	450.789	126. 399	129, 150	103.289	53. 193	115.911
				22, 156		16.731	16.116	10, 600	19, 305
	HEMBA1005002	73.560	26. 438		12.657				
	HENBA1006005	59.620	7.083	16.863	8.213	29.019	53.513	52. 130	23.838
	HEMBA1006011	25.811	30.413	39, 888	21.434	54.488	30. 978	27.996	25. 339
15	HEMBA1006013	51.604	13. 251	19.743	11.817	15.364	28.363	21.493	18. 574
								33, 305	26. 397
	HEMBA1006016	101.929	42.149	115, 996	36. 228	39.875	46.607		
	HEMBA1006019	31.772	18.482	22.979	15. 207	22.984	24. 244	26.246	14. 100
	HEMBA1006021	26.984	10.213	45. 937	9. 253	20.615	14. 587	14.203	12.296
		100.930	40.046	62.368	42.744	23.660	46.057	25.008	19. 323
	HEMBA1006022								
	HEMBA1006031	42.088	41. 281	14.729	11.264	12.725	36.716	13.037	5. 133
20	HEMBA1006035	10.089	10.059	27.290	8.123	6.309	6.629	2.019	5. 229
	HEMBA1006036	188.431	82.469	443.914	119.939	80, 135	81.126	54, 157	94.631
	HEMBA1006042	69.906	33.773	134, 462	30. 108	23.244	29.765	29.479	29.607
								2. 586	
	HEMBA1006044	53.721	10. 199	12.818	4.725	8.467	5. 436		4. 088
	HEMBA1006045	48, 078	43.730	61.128	28. 336	25, 311	26.461	23. 478	44.272
	HEMBA1006048	35.685	18. 435	41.495	19, 225	19,636	34.213	26. 302	28.809
			356. 500	78.844	24. 270	47.030	114.986	63. 574	385.970
25	HEMBA1006053	0.000						9, 079	5. 545
	HEMBA1006055	7.603	5. 331	12.625	4. 484	13.776	12. 227		
	HEMBA1006058	51.872	19.394	14.828	7.834	11.877	25.640	15. 830	21.486
	HEMBA1006063	72.886	52. 429	63.882	34. 021	30.125	39.536	28, 303	35.860
			14. 253	7. 505	3. 169	2.242	3. 352	4. 358	0.888
	HEMBA1006067	6.005							23, 976
	HEMBA1006081	70. 282	19, 151	25.838	8. 981	9,908	26. 560	16.837	
	HEMBA1006089	54. 392	23.145	42.709	18. 278	17.433	17.768	18. 372	23.981
30	HEMBA1006090	71.092	20.389	36.832	15. 386	17.868	38.904	35, 031	18.238
			28. 947	126. 425	16.353	30.302	56.034	53, 660	66.468
	HEMBA1006091	69.022							
	HEMBA1006093	111.885	11.435	50.738	16. 185	27.687	43.178	25. 048	14. 980
	HEMBA1006099	40.381	27.136	39.149	18. 199	31.100	31.158	28. 536	25.484
	HEMBA1006100	36.979	48.991	259. 267	41.090	50.094	24.833	13.379	34.466
			19.301	21.811	11.126	8.795	12.441	8,780	16.453
0.5	HEMBA1006108	40.170							
35	HEMBA1006114	42.849	44.783	46.702	33. 193	23. 220	34. 526	28. 294	51.756
	HENBA1006121	160.208	21.943	26.728	10.160	21.331	17.129	26, 838	25.137
	HENBA1006124	63.151	11.764	15.994	17.764	14.099	57.249	29. 200	8.240
	HEMBA1006125	72.730	70.405	57.020	50.057	45. 287	40.856	45. 865	68, 939
								25.726	21.315
	HEMBA1006130	36.221	31.688	34.742	7.817	28, 246	34. 473		
	HEMBA1006138	160.258	170.815	435. 120	106.719	139,660	100.947	67.854	89.604
40	HEMBA1006142	127. 194	85.725	238. 562	54.531	52.936	65.032	45. 938	59.791
40	HEMBA1006150	66.777	58. 231	76.666	59. 941	19.605	46.114	33, 261	75.731
				104. 921	29.648	46.546	66.736	74, 155	88. 383
	HEMBA1006151	189. 265	57.959						
	HEMBA1006155	141.288	19.560	50. 142	11.752	32.711	79.435	60.621	32.838
	HEMBA1006158	17.276	12.039	19.210	7.139	7.468	23.241	7.360	13.357
	HEMBA1006164	140.272	70.843	382. 965	97. 488	87.832	69.460	42.210	85. 135
					13. 911	21.700	40.783	26.049	37.233
45	HEMBA1006171	66.839	48.304	34.618					
	HEMBA1006173	63.939	35. 393	52.598	22. 894	32.403	35.413	40. 872	67.870
	HENRALOGS 176	51.671	222.661	52.703	39. 359	29.305	59. 271	24. 272	83.343
					29.730	26. 735	30.382	19, 907	34.405
	HEMBATOOS 176		38 362	1 (3/, 45)					
	HEMBA1006182	72.842	38.362	132.455					40 766
	HEMBA1006182 HEMBA1006197	72.842 16.655	38. 362 31. 338	37.528	55.808	23. 143	18.848	13. 456	40.765
	HEMBA1006182	72.842	38. 362 31. 338 15. 178	37.528 21.337	55. 808 16. 185	23. 143 25. 764	18.848 15.643	13. 456 14. 389	18.561
	HEMBA1006182 HEMBA1006197 HEMBA1006198	72.842 16.655 30.466	38. 362 31. 338	37.528 21.337	55.808	23. 143	18.848	13. 456	
. 50	HEMBA1006182 HEMBA1006197 HEMBA1006198 HEMBA1006213	72.842 16.655 30.466 38.783	38.362 31.338 15.178 20.120	37, 528 21, 337 38, 136	55.808 16.185 15.627	23. 143 25. 764 10. 604	18. 848 15. 643 25. 761	13. 456 14. 389 21. 716	18. 561 35. 282
. 50	HEMBA1006197 HEMBA1006198 HEMBA1006198 HEMBA1006213 HEMBA1006217	72.842 16.655 30.466 38.783 32.003	38. 362 31. 338 15. 178 20. 120 18. 510	37.528 21.337 38.136 33.960	55.808 16.185 15.627 4.079	23. 143 25. 764 10. 604 17. 107	18.848 15.643 25.761 31.016	13. 456 14. 389 21. 716 36. 526	18. 561 35. 282 19. 419
50	HEMBATOG5182 HEMBATOG5197 HEMBATOG5198 HEMBATOG5213 HEMBATOG5217 HEMBATOG5226	72.842 16.655 30.466 38.783 32.003 40.304	38, 362 31, 338 15, 178 20, 120 18, 510 60, 090	37. 528 21. 337 38. 136 33. 960 110. 529	55.808 16.185 15.627 4.079 40.359	23. 143 25. 764 10. 604 17. 107 39. 915	18.848 15.643 25.761 31.016 62.796	13. 456 14. 389 21. 716 36. 526 35. 202	18. 561 35. 282 19. 419 59. 281
. 50	HEMBA1006197 HEMBA1006198 HEMBA1006198 HEMBA1006213 HEMBA1006217	72.842 16.655 30.466 38.783 32.003 40.304 40.954	38. 362 31. 338 15. 178 20. 120 18. 510 60. 090 9. 021	37. 528 21. 337 38. 136 33. 960 110. 529 21. 361	55.808 16.185 15.627 4.079 40.359 7.280	23. 143 25. 764 10. 604 17. 107 39. 915 14. 241	18. 848 15. 643 25. 761 31. 016 62. 796 13. 056	13. 456 14. 389 21. 716 36. 526 35. 202 4. 951	18. 561 35. 282 19. 419 59. 281 7. 077
. 50	HEMBATODE 182 HEMBATODE 197 HEMBATODE 198 HEMBATODE 213 HEMBATODE 217 HEMBATODE 226 HEMBATODE 225	72.842 16.655 30.466 38.783 32.003 40.304 40.954	38, 362 31, 338 15, 178 20, 120 18, 510 60, 090	37. 528 21. 337 38. 136 33. 960 110. 529	55.808 16.185 15.627 4.079 40.359 7.280	23. 143 25. 764 10. 604 17. 107 39. 915	18.848 15.643 25.761 31.016 62.796	13. 456 14. 389 21. 716 36. 526 35. 202	18. 561 35. 282 19. 419 59. 281
50	HEMBATOG5182 HEMBATOG5197 HEMBATOG5198 HEMBATOG5213 HEMBATOG5217 HEMBATOG5226	72.842 16.655 30.466 38.783 32.003 40.304	38. 362 31. 338 15. 178 20. 120 18. 510 60. 090 9. 021	37. 528 21. 337 38. 136 33. 960 110. 529 21. 361	55.808 16.185 15.627 4.079 40.359 7.280 10.747	23. 143 25. 764 10. 604 17. 107 39. 915 14. 241	18. 848 15. 643 25. 761 31. 016 62. 796 13. 056	13. 456 14. 389 21. 716 36. 526 35. 202 4. 951	18. 561 35. 282 19. 419 59. 281 7. 077

445

Table 36

									·
	HEMBA1006252	36.069	24.612	74. 170	29.506	28.055	19.517	14.085	15. 356
	HEMBA1006253			20.773	16.455	11.705	12.936	6.506	11.398
		75.854	7.002						
_	HEMBA1006259	37.456	48.402	136.000	39.735	19.462	25. 242	19.832	25. 931
5	HEMBA1006261	23.677	23. 578	5. 874	13.012	7. 127	69.427	7.141	17. 143
				30.879	8.970	7.077	19.793	22.288	18. 289
	HEMBA1006268	35.886	12.563						
	HEMBA1006271	122.980	98.618	185. 469	77.610	45.268	47.910	36.533	47. 867
	HEMBA1006272	16.261	12.829	9. 415	4.968	5. 925	27.765	15. 997	7.567
						30.445		18. 419	22. 459
	HENBA1006273	47.890	12.641	71.219	20.880		30. 473		
	HEMBA1006276	79.296	11.878	30.854	34.032	8.760	27, 168	16. 165	7. 501
10		40.093	7.717	26.091	4.506	18.669	11.680	11. 224	9.893
	HEM8A1006278							28. 267	
	HEMBA1006283	16.994	23.586	25. 614	25.226	23. 447	59.086		25. 848
	HEMBA1006284	29. 982	22.166	27. 891	20.874	8. 594	18.386	15, 293	13. 396
			13.071	36.861	9.670	4.059	11.649	31.851	7. 519
	HEMBA1006291	22.745							
	HEMBA1006292	17.718	8.916	20.081	10.169	4. 378	7.903	9. 259	7.898
	HEMBA1006293	31.307	10.056	8.749	4.645	4.097	6,631	8.473	7. 189
15			5.917	6. 157	1.371	4. 543	2.465	1.701	2.648
7.5	HEMBA1006299	21.091							
	HENBA1006309	69.975	25.568	110.869	33, 191	19.510	31.160	24.850	17.764
	HEMBA1006310	40.983	23. 265	36. 585	20.570	11.748	29.056	27. 263	17. 748
				64, 711	8.925	20.171	92.798	9.481	19. 313
	HEMBA 1006311	85.398	20.844						
	HEMBA1006313	27.762	12.975	47.707	17.417	7.455	13. 117	9.891	6. 082
	HEMBA1006316	23.345	3.751	3. 303	2.158	8.774	9.668	8.505	3. 270
				185. 981	41.111	28.820	37.527	35. 377	85. 968
20	HEMBA1006328	79.937	83.744						
	HEMBA1006334	22.524	16.717	17. 679	5.994	8. 506	9.813	3.866	5, 361
	HEMBA1006335	72.666	41.477	35, 235	27.435	6.110	5.851	24. 375	8.434
				132.978	45.518	34.812	40.158	41.934	25. 330
	HEMBA 1006344	34.707	67.866						
	HEMBA1006347	34.301	16.445	32. 190	19.603	16.749	20.762	20.884	15. 376
	HEMBA1006349	139.389	26.300	48. 767	43.275	22.026	24.648	22.876	21. 499
			17.873	15. 526	9,410	8.472	14.845	7.491	9.414
<i>25</i>	HEMBA1006352	21.127							
	HEMBA1006357	94. 337	82.319	287. 531	67.888	76.120	47.179	41.500	59. 557
	HEMBA 1006358	48. 925	31, 345	132. 494	32.473	25.019	28, 197	13. 250	24. 899
	HEMBA1005359	57. 203	18. 522	160. 314	70.923	17. 441	30.686	11.154	47. 991
							13.876	6.804	
	HEMBA1005360	29.518	10.133	15. 515	17.275	6.141			8, 351
	HEMBA1006364	59.236	7. 900	27. 522	12.114	5. 401	15. 432	17, 981	6. 672
	HEMBA1006377	67.120	31, 113	57.269	33.567	23.849	45, 246	31.609	20. 280
30		30.120					33.416	23.616	40. 932
	HEMBA 1006380	73.227	57.029	182. 581	57.870	22. 288			
	HEMBA 1006381	359.346	122. 755	376.090	126.304	112.826	145.346	91.469	93. 252
•	HEMBA1006385	60.234	62.166	257. 945	59.429	59. 157	40.136	35. 385	17. 281
						16. 455	41.253	16.013	27.609
	HEMBA 1006390	71.393	38.752	46.828	25.848				
	HEMBA1006391	61.261	18.765	20. 686	10.972	10.022	39.431	27. 305	11.797
	HEMBA1006398	42.089	3.225	18. 036	5.299	25.386	· 6. 480	0.000	3. 308
35				40. 904	17.896	18. 180	84. 926	41.325	24.773
33	HEMBA1006405	137.413	28.645						
	HEMBA1006410	149.580	32.840	61.022	20.027	39.718	54. 551	23.826	33. 928
	HEMBA1006416	96.031	62.892	198.896	50.538	38.551	37.025	37.809	33.447
	HEMBA1006418	23.236	18. 335	23.851	11.378	10.280	28, 208	46.245	36. 223
								40.042	52. 384
	HEMBA1006419	189. 293	101.979	476. 145	90.626	79, 213	64. 306		
	HEMBA1006421	39.702	26. 487	127. 221	23.773	16.184	14.450	12. 270	13. 523
40	HEMBA1006424	4. 484	36. 452	10. 588	3.778	4. 512	7.346	2. 324	3. 323
40			67. 224	230.530	60.836	32.273	40.489	17. 284	36. 244
	HEMBA 1006 426	88.597							
	HEMBA1006430	61.672	17.989	69. 151	15.913	11.038	15.595	9.696	17. 632
	HEMBA1006438	45.084	34.475	111.512	27.012	15.035	34.111	12.678	11.056
					9. 326	15.672	34.167	27.442	18. 331
	HEMBA1006445	48. 245	13.919	53. 981					
	HEMBA1006446	22. 911	3.160	3. 324	1.568	4. 341	2.585	1.331	0.000
	HEMBA1006456	36, 915	28. 165	141.114	1 18. 927	65.823	33.549	13,651	33. 405
45				161.108	40.447	22.274	32.823	18.018	27. 165
	HEMBA1006461	60, 747	42. 392				+		4 555
	HEMBA 1005467	13.357	6.130	15.734	10.759	4.032	4.4/1	6. 183	2.655
	HEMBA 1005470	73, 950	30.706	103, 625	27.235	29.870	33.756	33.818	24. 286
			4. 504	7. 503	2.933	2. 522	5. 224	10.020	1.873
	HEMBA1006471	19.032							
	HEMBA 1006474	25.718	12. 420	21.381	11.498	9.614	19.875	17.655	13. 491
	HEMBA1005476	180.042	91. 936	63.588	43.462	42.248	109.725	88.725	65. 945
50	HEMBA1006482	129.627	169. 312	167. 982	151.338	57.839	95. 521	75. 480	239. 325
	HEMBA1006483	99.620	64.773	232.207	50, 445	29.074	37.572	23, 818	27. 130
	HEMBA1006485	41.690	4. 055	17.445	11.682	4. 522	9. 351	6.411	10.066
	HEMBA1006486	76. 250	36. 421	29.634	46.687	17.302	21.229	17.832	15.706
	HEMBA1006489	5.771	32.673	2.141	5. 240	2.356	4. 324	4.739	7. 328

446

Table 37

	HEMBA1006492	14.002	19.916	24.662	35. 451	8.836	8.075	11, 419	12.090
								5. 851	5. 887
	HEMBA1006494	7.279	0.000	19, 790	3.750	8.718	8. 343		
·	HEMBA1006497	41.284	12.396	23. 326	6. 590	7.186	11. 228	9.062	5. 781
5	HEMBA1006501	160.565	16, 895	26.893	13, 446	17.608	65, 467	41.560	6. 197
						10.919	15.054	17. 536	15.658
	HERBA1006502	53.451	19.114	39, 593	25. 366				
	HEMBA1006507	19. 274	8, 180	10.287	4. 521	7.939	5. 288	15.480	10.062
			30.085	91.871	18, 732	21.918	45.881	29.819	16.672
	HEMBA1006517	95. 989				9.774	14. 205	14.646	13. 907
	HEMBA1006521	31.224	27.873	37.864	18. 318				
	HEMBA1006529	28.702	20.010	34.050	20. 150	16.588	7.353	8.993	17. 327
10		18, 445	16.411	29.175	14. 433	12.214	15.734	15. 731	8. 081
	HEMBA1006530					8, 160	19.824	5. 837	3. 457
	HEMBA1006535	11.627	7. 208	18.048	3.956				
	HEMBA1006536	68.087	40.009	142.475	43. 263	34. 343	42.050	42. 157	23.975
	HEMBA1006540	20.393	10.867	35. 153	8.637	8.656	15.027	11.094	10.350
				59, 940	7. 791	7.169	15.883	8.745	8.693
	HEMBA1006544	30, 281	4, 662					22. 328	34.045
	HEMBA1006546	68.722	53. 155	127. 193	49. 337	73.807	60. 506		
15	HEMBA1006549	13.885	13.666	21.800	11.666	8. 491	14. 211	8. 987	6.080
	HEMBA1006559		22.040	38, 197	16.910	14.550	14.058	13.018	17. 217
		26.976					26.651	18.158	19, 510
	HEMBA1006562	55.924	24.663	75.789	20. 363	17. 181			
	HEMBA1006566	20.849	6.116,	14. 933	8.767	9. 572	6.937	5. 229	4. 788
	HEMBA1006569	67.508	20. 299	44, 291	27.048	12.798	15. 243	24. 739	31.861
				15.862	1.796	3. 407	11.582	8.381	5, 922
	HEMBA 1006572	21.817	4. 339						5, 110
20	HENBA1006579	5. 427	18. 336	4.219	3. 440	2.139	5.460	3.967	
	HEMBA1006583	31.967	15.854	29.307	14.271	11.747	26.889	17.058	10.451
	HEMBA1006595	59.014	41, 577	148.359	30.660	16.681	19.571	13. 265	24.768
				210, 001	47.574	27. 392	47.009	27.887	28.666
	HEMBA1006597	111.817	64. 480						31. 587
	HEMBA1006606	79.184	47.311	131.822	40.177	33. 228	35, 403	25. 240	
	HEMBA1006612	43, 105	20. 909	46, 913	39. 205	20.348	25. 383	18.706	17.150
	HEMBA1006617	79.139	62.924	235. 236	60. 258	30.407	40.264	28. 184	38.643
25							291, 427	208. 533	65.478
	HEMBA1006624	449.384	84.050	165. 494	39.352	209. 908			
	HEMBA1006631	168.309	108.316	381,778	89.696	71.812	80.634	39. 325	50, 996
	HEMBA1006635	51,406	33.730	158, 286	28.605	19.347	19.781	9.639	12.894
			30. 354	51.867	15. 409	33. 210	43, 083	25.295	12.985
	HEMBA1006639	67.363					68.642	41.724	17, 931
	HEMBA1006643	229.685	30. 246	56.218	16. 406	35.196			
	HENBA1006648	80.985	32.464	39, 607	14. 926	36.718	12.135	32.217	48, 853
30	HEMBA1006652	118, 455	69. 232	231.917	50.609	51.023	50.716	21.698	29. 527
				46.472	16. 579	12.358	15.364	13.867	9. 224
	HEMBA1006653	46.971	16.614					33.470	16.872
	HEMBA1006658	89.823	28. 353	60.976	37.660	28.124	47.014		
	HEMBA1006659	79.863	33.626	48. 217	49. 132	29. 124	33.070	25. 182	33.784
	HEMBA1006665	25.726	26.740	39,661	13.975	13. 287	15. 240	12.046	10.419
					6.319	4. 257	10.392	2.791	2, 171
	HEMBA1006666	8. 276	4. 281	10.565					
35	HEMBA1006671	39. 553	178.623	135. 413	18, 941	17. 294	37.782	10. 166	32.048
	HEMBA1006674	100, 472	44. 108	175.724	46. 922	36. 367	44. 809	43.576	43.269
	HEMBA1006676	120.417	42.888	163,816	29.504	40, 435	60.162	32.540	34.825
					4.035	8. 982	19.092	3. 958	0.000
	HEMBA1006682	27.104	2.556	23.174					
	HEMBA1006688	57.351	56.288	111.358	60. 597	65. 322	37. 545	20. 757	20.789
	HEMBA1006695	132.496	140.334	315.655	97. 296	56. 206	54.392	37.622	57.596
	HEMBA1005696	65.136	25. 204	42, 137	26.654	26. 490	30.156	6. 159	27.512
40				8.881	7.114	3. 362	1.846	7,796	1.965
	HEMBA1006702	4.275	4. 328						
	HEMBA1006707	52.417	20.766	26,862	21.409	19.843	32.229	13. 146	18.546
	HEMBA1006708	126.875	38. 520	66.803	31.253	33. 294	55.347	32.071	18.229
	HEMBA1006709	67.500	31.686	94. 432	24. 924	17. 365	30.329	18. 503	23.474
					12.664	16.091	54.326	26.752	11,544
	HEMBA1006717	110.641	21.536	29. 255					17.797
45	HEMBA1005724	34. 421	23.073	25.607	18. 231	12.305	27.570	10.585	
45	HEMBA1005731	36, 072	18. 255	41.441	15. 382	16.479	17.272	10.826	15. 482
				30.096	14.542	20, 232	22.606	10.316	11,440
	HEMBA1006737	60.467	14, 107					15. 223	23.529
	HEMBA1006742	60. 258	45. 190	134.964		21.315	21.889		
	HEMBA1006743	41,970	22.864	31.760	22.024	15, 126	23.989	13.179	16.281
	HEMBA1006744	181.058	97.273	433,004		69.785	59.354	46,770	61.806
							34.516	28. 426	23.238
50	HEMBA1006749	51.776	9.753	37.994		23.164			
50	HEMBA1006752	124. 800	50.318	88.111	59.765	47.490	69.451	37.541	47.074
	HEMBA1005754	49. 957	30.459	86.726	23.747	17.745	16. 269	10.783	12.424
			21.737	26. 190		18.249	38. 492	30.654	15, 933
	HEMBA1006758	75, 460							8, 573
	HEMBA1006767	14. 002	15. 106	11.961		5, 628	13.334	8. 382	
	HEMBA1006770	120. 485	21.505	62.144	29.559	32.512	49,739	45. 952	28.318
					<del></del>				

447

Table 38

	HEMBA1006779	81, 492	51.077	162.657	41.153	39, 166	36.722	18.025	29.256
	HEMBA1006780	78, 359	78.052	345.442	73.371	68.858	55. 888	41.524	39.494 -
			21.233	20.440	14. 349	11.547	38, 549	19.736	25, 701
5	HEMBA1006789	29.455			55.068	49.500	46. 284	21.141	40.750
	HEMBA1006795	143.727	88.701	218.732				15. 525	15. 352
	HEMBA 1006796	87.214	15.814	115. 542	17. 685	16.790	38.694		30. 733
	HEMBA 1006805	68.116	31.212	153.041	33. 162	30. 301	34. 197	24. 275	
	HEMBA1006807	94, 524	86.723	157.559	64. 349	36. 505	62.933	23.097	55. 508
	HEMBA1006813	40, 696	4, 415	4, 750	4. 264	10.978	7.562	6.201	3. 198
	HEMBA1006819	53,717	15.217	30.071	14.679	17.006	30.866	20. 346	6. 250
10		39.052	30. 425	111.325	35.769	34. 975	22. 216	18. 924	20.698
	HEMBA1006821			201.721	47. 107	40.322	27. 255	21.689	27.074
	HEMBA1006824	68.491	61.498			40.147	75. 996	66.799	71,706
	HEXBA1006832	84.462	89.500	102.038	77.046	41, 460	61, 443	30. 402	31. 940
	HEMBA1006834	123.958	57.085	160.407	48. 909			22, 426	16. 742
	HEMBA 1006835	33,705	19. 529	38. 470	23, 193	18, 979	22.344		
	HEMBA1006843	52.436	44. 642	96.773	258.615	195.878	33.141	8.256	13. 117
15	HEMBA1006849	88, 931	34.224	158.388	39.483	30. 349	34. 943	15.743	28. 240
	HEMBA1006850	44.733	24, 923	67.667	24. 186	15.829	36.593	11.223	18. 454
	HEMBA1006861	215. 207	94. 180	158, 997	67.349	259. 512	135.856	371.932	44.063
	HEMBA1006865	124.996	59.773,	124. 376	43. 328	59.356	71.072	66.350	45. 129
			11.094	39.646	14. 084	12.902	11.855	5. 865	18.338
	HEMBA1006867	16.632			5.013	6. 262	5. 127	7. 141	8. 422
	HEMBA1006873	9. 965	9. 279	7.010			13. 165	16. 493	9. 073
20	HEMBA1006877	44.043	18. 321	20.546	8. 172	14.670		41.513	17. 905
	HEMBA1006878	100, 427	34, 418	109.029	25.739	29. 525	48.800		
	HEMBA1006879	108. 299	42.811	121.051	60.872	47.507	40.075	14. 429	51. 924
	HEMBA1006884	95, 426	29, 331	67.556	27.787	25.909	106.818	47.878	47. 793
	HEMBA1006885	107, 720	54. 342	127. 920	62.272	55. 739	51.739	36.790	50.612
	HEMBA1006886	50.841	22, 970	51, 528	12.561	20.660	23. 207	26.952	19, 149
or	HEMBA1006889	81.809	20. 952	21.474	12.691	24.681	41.822	48.768	15. 196
25	HEMBA1006896	68.030	97, 285	75.370	52.746	23. 109	44. 481	37.701	50.662
			36,410	61.016	23.329	21, 390	38.404	27.583	22.774
	HEMBA1006900	61. 515	19,713	47, 129	12.105	11.602	27.830	26.548	13.885
	HEMBA1006902	43.283			78. 230	79.588	63.729	39.994	64. 953
	HEMBA1006912	183.904	90.995	338. 160		25.895	38. 586	22.479	33.810
	HEMBA1006914	54. 548	39.053	48. 945	35.736				30.818
30	HEMBA1006916	62.872	0.000	65. 115	29. 982	32.625	61.537	62.750	
30	HEMBA1006921	64.867	21.840	74.902	15.692	30.866	41. 257	25.569	10.362
	HEMBA1006926	51. 195	10.616	76, 671	24. 435	20.300	84. 402	29.503	20.967
	HEMBA1006927	24.016	13.778	23. 573	5. 335	15.250	11. 291	11.672	7.086
	HEMBA1006929	7. 146	8.487	5. 431	5. 526	1.676	5. 970	5.688	3. 134
	HEMBA1006936	68. 233	22.847	45. 566	20.391	16.346	25. 493	20. 196	17.720
	HEMBA1006938	14. 202	8, 409	31.234	7.743	5. 002	6.780	6.773	5. 945
35	HEMBA1006941	30. 559	24.290	40.928	13.779	16.040	34. 253	22.542	18.507
	HEMBA1006942	147.487	57.842	121.883	69.207	55. 456	76.853	61,942	65.640
		80.545	64. 930	104.037	63.709	40.444	54.676	33.533	31.915
	HEMBA1006945			23.921	1.860	15.813	7. 071	10.866	5. 231
	HEMBA1006949	10. 292	41.467			18. 283	39.764	15. 332	12.456
	HEMBA1006952	58. 685	12.572	34.750	8.032		36. 160	36.715	34. 791
	HEMBA1006960	91. 939	38.895	93.164	24.834	34.400			
40	HEMBA 1006 973	74. 208	24. 793	50, 621	17.619	22.844	24. 971	24.844	16. 167
	HEMBA1006974	48.691	39.013	59.414	48.064	16.799	38. 579	21.301	46.006
	HEMBA1006976	35. 907	15.675	32.116	19.091	14. 522	30.574	25.042	18.348
	HEMBA1006989	6, 422	2.207	2.374	3.336	2.670	3.696	2. 557	3. 536
	HEMBA1006993	334, 266	64, 150	357.947	46.138	95.466	144.777	109.174	54.000
	HEMBA1006996	9, 183	9.870		9.483	5.722	9.518	8.368	9.637
	HEMBA1007001	117.610	95, 668		56,093	55. 288	47.863	27.205	56.828
45			41.846		21.453	16. 249	59.722	46.434	40.628
	HEMBA1007002	93. 134				20. 783	34. 293	29, 163	29. 338
	HEMBA1007013	65.734	23.106				17. 982	16.658	15.035
	HEMBA1007016	36.649	14.972						
	HEMBA1007017	6.290	0.000				2. 329	1.949	0.000
	HEMBA1007018	19.457	15.664				15.084	9.105	14. 124
	HEMBA1007044	139.784	50.078	125.738	15.913		123. 367	90.838	36.173
50	HEMBA1007045	49.576	7.913		9.069	10.104	19.099	12.683	7.276
	HEMBA1007051	36.374	44, 117				24.088	15.546	9. 363
	HEMBA1007052	69. 582	19.611				19.409	18, 969	10.939
		25, 326	27.611				20.128	7.847	9. 544
	HEMBA1007053						36.241	14.769	
	HEMBA1007057	45. 897	13, 545	33.857	18.616	25. 861	1 30.241	14. (09	1 13. 302

448

Table 39

			• •	14	016 33				
	HEMBA1007062	129.012	18.903	40.670	21.323	29, 469	40, 252	29.408	1 15,700
	HEMBA1007063	81.681	45. 884	187.380	52. 391	36.943			
	HEMBA1007066	98.395					28.608	35.303	41.236
5			32.970	35.373	22.961	11.085	42.430	26.631	14.760
_	HEMBA1007069	23. 449	21.519	78.409	16.835	27. 425	17.217	9.095	16.163
	HEMBA1007073	54.833	42.548	40.682	29. 352	11.879	7.937	24. 282	19.372
	HEMBA1007076	83.020	48.746	248.260	61.189	50, 193	68.045	43.836	35, 550
	HEMBA1007078	151.561	159.600	446.445	189.146	130. 283	98.734	65.934	117,079
	HEMBA1007080	43.963	44,765	174.545	66.950	45.879	43.194	43.909	50, 100
	HEMBA1007084	78, 948	60,672	268. 327	63.769	63.088	60. 307	35.006	46.866
10	HEMBA1007085	263.538	108.018	162.599	48.155	77. 545	161.321	63.614	80.640
	HEMBA1007087	85. 598	25, 085	47.862	25. 580	13, 918	62.815	143.461	
	HEMBA1007089	21, 131	32.023	21.145					30.856
	HEMBA1007095				14.738	7.213	19.681	9.036	10.026
		147.777	215.051	136.910	63.992	170.706	117.992	103.152	86.452
	HEMBA1007101	78.959	53.790	147.891	35.676	28. 082	27. 200	19.131	25. 922
15	HEMBA1007104	66.308	23. 279	45.417	11.902	19, 468	48.054	26.760	16. 647
	HEMBA1007106	28. 449	17.761	41.268	28.670	17.681	14, 174	10.999	7.534
	HEMBA1007112	12.759	8.412	16.340	9.319	7, 661	7.304	13.296	6.622
	HEMBA1007113	126.702	0.000	229.408	64.551	40. 242	39.032	13.319	26.174
	HEMBA1007121	219.036	207. 410,	696.658	149.217	168. 827	131.628	642.099	128.755
	HEMBA1007129	50. 726	42.510	63.847	31.663	26. 417	24.371	18.928	20, 103
	HEMBA1007147	111.299	117.722	312.811	79.949	67. 395	74.391	35.758	
20	HEMBA1007149	83, 453	6.442	19.831	7. 332	11.043			54. 184
	HEMBA1007151	97, 211	33. 530	53. 944			9. 349	9.831	8.756
	HEMBA1007172	52.683			24. 544	18.501	35. 246	36.228	24. 174
	HEMBA1007174	52. 921	25. 324	438.704	42.182	28. 599	38. 126	26. 167	25.770
			13. 482	44.770	21.384	19.520	28. 559	22.332	20.471
	HEMBA1007176	89. 919	24. 768	53, 414	32.841	44.643	73.679	87.040	30.762
25	HEMBA1007178	93. 941	73. 120	135. 427	34, 313	32.040	34.622	22.898	24.897
23	HEMBA1007185	62. 558	18, 807	36.824	15. 490	20. 528	37.568	22.260	12.783
	HEMBA1007186	70.967	31.546	59. 038	21.059	21, 332	35.648	42.864	11.346
	HEMBA1007194	53. 376	38, 911	126.660	33.992	23.875	21.109	12.122	23, 307
	HEMBA1007200	74. 955	53.829	44. 212	23.979	20. 225	32.762	55, 417	22.176
	HEMBA1007203	87.803	26.807	41.357	14.648	9. 791	23.392	30, 167	17.274
	HEMBA1007206	82,800	73.675	225. 293	44. 451	28.674	37.091	14,673	34, 505
30	HEMBA1007224	25. 614	40. 402	50, 116	21.484	14. 920	22.548	13, 197	20.053
	HEMBA1007226	88, 512	43.606	93, 121	22. 209	17.911	38.704	43.759	31.721
	HEMBA1007240	131.657	62.804	86,650	9. 510	21.890	53.116	42.250	16.655
•	HEMBA1007241	12.225	7.719	18.461	5. 051	6.724	15.945	3.135	
	HEMBA1007242	21, 409	14.030	13.648	11.068	6. 265			5. 390
	HEMBA1007243	61.824	25, 854	40, 264	17, 235		17. 370	8, 487	5. 236
35	HEMBA1007251	37.660	16.946	37. 149		23.438	39. 197	31.904	20. 347
	HEMBA1007256	53, 905	43.642		16.699	12.180	19.482	30.321	10. 262
	HEMBA1007267	80.741		113, 110	31.642	27.946	30.492	18.548	23. 645
	HEMBA1007273		40.085	207.160	61.174	38. 220	29.008	32.292	29.672
	HEMBA1007279	41.062	9.087	11.906	5, 193	6.445	7.723	9. 225	4. 483
		54. 376	20.734	133. 494	27. 987	21.355	19.941	17.364	19. 503
40	HEMBA1007281	8. 523	5.717	4,731	3. 403	2. 317	2.497	2. 740	0.000
40	HEMBA1007283	25. 940	14.444	24. 974	23. 487	19.771	23.418	19.378	26.409
	HEMBA1007288	57, 959	39.576	155. 227	28.725	24. 689	25.110	16.998	16.095
	HEMBA1007291	37.974	19.069	59. 253	20.445	13.404	17.376	13.060	13. 147
	HEMBA1007299	446.640	93.668	199.852	61.423	94. 129	249.345	241.373	85. 323
	HEM8A1007300	103.752	25.694	24, 914	18.217	40.413	26.018	31.407	16.669
	HEMBA1007301	49.752	18.178	32.677	18.170	33.650	33.786	22.892	12. 782
45	HEMBA1007319	13.312	10.598	23. 453		4, 278	9, 382	2.996	8.570
	HENBA1007320	53,723	23, 595	62.301	29.439	16.672	32.932	28. 191	18. 418
	HEMBA1007322	45. 986	125.362	77. 545	43.693	17. 955			
	HEMBA1007323	64.720	16.869	22.970			45.689	39.556	80. 836
	HEMBA1007326	313.094	189. 188		11. 238	11.687	32.209	25. 350	7. 506
	HEMBA1007327	78.767		862.276	214.045	178.109	171.587	70.819	115. 174
50	HEMBA1007332		61.102	219.980	55.002	29, 411	44.095	29. 354	42. 286
50		71.516	9.318	34.879	5. 559	7.452	24.826	12.763	20.050
	HEMBA1007341	89.805	53. 431	207. 395	82.402	105. 877	47.861	32.826	50.162
	HEMBA1007342	22.063	17.289	28. 253	18.196	17.751	26.378	13.820	9. 173
	HEMBA1007347	112.392	64. 499	230.022	60.348	47.557	83.758	30.683	33. 285
	HEMBA1007353	1.685	3. 520	0, 575	1.850	1.976	0.107	2.081	0.788
	HEM881000005	60.047	46.027	121.870	38.241	20.699	18.268	20.068	26.957
55	HEM881000008	97. 929	53, 604	274. 179	68.681	38. 935	39. 328	26.881	34. 873

## Table 40

	COPUSSA SA	744 744 7	*** ***	200 100 1	20 410	57.867	AC 202	75. 902	92. 924
	HEMBB1000018	122.130			20.419		95.203		
	HEMB81000024	181.606	97.019		02. 401	70. 406	70.591	40.304	66.798
5	HEMB81000025	85.919	29.049	45.055	23.789	13.946	24. 397	29.349	13.072
3	HEMBB1000030	108.167	68.316	303.677	83.010	68. 378	81.687	34.886	37.617
	HEMBB1000036	107.960	11,573	50.484	11.277	20. 480	41.381	25.378	14.730
	HEMBB1000037	77.688	29.380	69.658	56. 679	27.020	54.052	30.086	15. 311
	HEMB81000039	52.550	48, 503	140.795	30.096	18, 739	26.012	15, 151	21, 723
	HEMBB1000044	134, 136	75. 469	218.667	61.596	32.667	29.659	43.360	42.831
				31.004	18. 291	11. 321	20, 120	21.506	15.078
10	HEMB81000048	17.937	21.052				25. 584	18.572	17. 494
	HEMBB1000050	74.210	33.681	207.484	35, 691	22. 905			
	HEMB81000054	68.273	47. 191	245.350	44.008	24. 522	29. 259	22.570	21. 316
	HEMBB1000055	72.875	112.284		110.297	21.358	70,636	93.824	132.288
	HEMB81000059	331.577	184.587	662.540	182.481	130.065	131.364	90.002	121.903
	HEMB31000072	240.733	98.890	326.893	75. 919	67.742	118.222	108.108	91. 458
. =	HEM881000081	23.738	27.174	85.100	21.146	30.856	20. 458	7.513	15. 351
15	HEMB81000083	120.759	58. 163	188. 224	40.609	37.789	59, 334	33.712	39. 101
	HEMB81000089	67,618	54. 952	191.832	56.629	24.609	36.847	30.680	26. 912
	HEMB81000034		116.828	161. 958	31.504	29.300	49.613	36.239	35. 197
		355. 534				22.834	34.068	18.547	16. 455
	HEMBB1000097	27.834	63.724	51.488	14. 249				
	HEM8B1000099	157.641	91.912	456.470	71.078	50.739	64. 471	32.108	43.354
20	HEMBB1000103	75.781	59. 392	114.974	44.216	31.915	47.628	23.669	56. 268
	HEMB81000106	62.814	44. 996	77.918	35.044	19.825	40.409	26.156	46.001
	HEMBB1000113	43.660	33. 435	95. 987	42.744	. 19.714	20.114	15.899	21.606
	HEMBB1000119	57.350	21.211	42.528	17.770	19.517	28.754	23.570	30.104
	HEMBB1000133	92.950	65.230	58.619	69.544	53.706	104. 229	39.058	80.858
	HEMB81000134	44, 120	20. 554	76.693	40.611	24.712	37.185	42.327	21.963
	HEMB81000136	21.810	7.191	44.517	15, 599	7.339	22.582	12.399	24.899
25	HEMBB1000141	163.867	99.946	331.822	95, 807	55.858	64.560	36.737	52.602
	HEMBB1000144	96.831	97.019	183.423	88. 529	36.185	15, 577	29.259	32.144
	HEMBB1000147	59. 253	9.088	62.426	7. 391	11.451	7. 175	11.502	10.693
			28.723	34. 597	15, 309	19. 424	32.469	29. 105	19, 117
	HEMBB1000152	56.391					27.298	20, 174	15. 366
	HEMBB1000154	85. 308	47.878	101.061	33.881	19.477			
20	HEMBB1000155	35,691	35. 132	109.038	28. 164	29.608	22. 283	16.557	17.041
30	HEMBB1000173	170.611	173.001	494. 253	143.666	83.705	123, 932	65.317	76. 388
	HEMBB1000175	32.273	19.114	23.481	10.948	4.039	29.180	7.135	13. 322
	HEMB81000176	56.984	51.334	90.749	69.004	40.144	52. 980	25.845	19.359
	HEMBB1000198	70.426	12.768	26.381	10.237	6.266	11.215	8.858	5. 363
	HEMBB1000208	42.474	8.966	34.929	10.418	12.883	9.285	12.335	7. 978
	HEMB81000209	43.846	10.700	9.943	10.934	8.858	12.135	9.049	4. 168
35	HEMB81000212	27.532	12.579	76.077	15.361	33.518	17.471	13.132	16. 552
	HEM8B1000215	178. 324	89.053	294.606	95.420	68.598	89.720	51.270	61. 235
	HEMBB1000217	148.073	45.416	96.614	47.569	37.572	89.989	48.073	33.510
	HEMB81000218	88.298	123.000	347.859	84. 124	41.828	57.417	21.147	34. 605
	HEMBB1000226	70.693	14.949	41.586	31.786	30. 261	28.577	14.779	27, 177
	HEMBB1000230	28. 681	8.910	13.549	5.500	3.547	9.616	6.632	3. 293
					10.455	4. 589	41.554	8. 171	7.082
40	HEM881000240	44.662	12.588	13.211	18.503	18.758	11. 192	2.111	13.188
	HEMBB1000244	22. 390	13.510	42.652				9.540	2.708
	HEMBB1000250	20.878	6.254	20.741	9.109	1.841	13.561		
	HEMB81000258	101.717	75.034	336. 781	79. 281	52, 103	67. 231	33.313	34.880
	HEMB81000254	99. 327	57.280	269.540	83.791	39.799	96.654	62, 346	79.783
	HEMBB1000266	70.747	23.082	23.217	14. 455	28.745	34. 547	15.022	15. 672
45	HEMBB1000272	14.990	14. 502	10.270	6.954	12.730	6.133	4.205	16.611
	HEMBB1000274	105, 245	46.925	190.978	49.759	41.568	43. 127	18.199	25. 826
	HEMBB1000276	6.479	2.218	2.501	4.783	1.754	2.070	2.079	1. 252
	HEMBB1000284	4.790	5.088	7.884	3.489	2.213	3.213	1. 981	3, 304
	HEMB81000307	52. 330	30, 191	128.450	28.961	22.039	15.869	9, 113	21.677
	HEMBB1000309	85.347	36. 463	96.140	43.964	34.442	33.118	18.805	21.507
	HEMB81000312	41.862	30. 986	40. 349	24. 933	7.383	79.360	24, 114	16.788
50									
	HEM881000317	49, 311	18.053	26. 189	10.490	10.102	21.107	12.632	13.384
	HEN881000318	87. 180	33.847	208. 954	43.556	23.043	27.764	9, 191	17.641
	HEMB81000332	3.892	11.256	14.087	42.331	28.145	14. 132	2, 408	14.319
	HEMBB1000335	27.939	30.864	21, 167	28.071	12.651	30.027	12.746	21.753
	HEMBB1000336	68.463	26.023	48.843	10,608	22.871	23.654	23.868	13.927
55	HEM881000337	289.853	59.290	93. 527	52.168	54. 197	125.769	126. 562	60.614
				*					

Table 41

	G.W. D. C. A. A. A. A. T.		72 342 1	144 166 1	44 614 1	17 724 1	40 665 1	17 700 T	29. 721
	HEMBB1000338	54.685	45.765	123.480	44.612	17.722	26.663	17.708	
	HEMBB1000339	144.258	108. 124	265. 125	105. 421	89.798	89.055	55. 944	50. 241
5	HEMBB1000341	113.271	45. 522	132.906	32.751	40.166	37.986	28.017	30.881
-	HEMBB1000343	130.737	71, 935	259.845	80. 183	46.681	45.761	43. 928	46.721
	HEMBB1000354	202.146	151.264	495.642	157. 908	153.529	142, 579	67. 161	105. 322
	HEMB81000358	92, 244	22.827	29.160	24.670	22.387	48.989	59. 506	28.803
				97.758	27.483	21.576	23.750	17.278	16.569
	HEMBB1000369	55.720	25.874			38, 792		31.504	45. 653
•	HEMBB1000373	52.572	59, 105	70.779	61.379		44. 185		
10	HEMBB1000374	153. 545	115.183	389_274	108. 150	98.073	80.319	58.214	75. 906
	HEMBB1000376	95. 394	132.554	369.986	146.818	60.328	63.876	73.647	43. 202
	HEMBB1000383	37.023	35, 429	24.954	13.017	10.381	22.638	16.842	8.781
	HEMB81000391	127. 327	30, 055	105. 971	24.962	30.891	57.827	37.484	11.921
		35, 143	10.865	22.406	8.561	4, 100	8.569	2.643	8.889
	HEMB81000399			44.945	25. 430	13.012	19.024	7.725	18.695
	HEMBB1000402	82.616	20. 485		8.593	9. 455	9. 301	2.672	7. 956
15	HEMBB1000404	18.903	12.568	10.300					3. 407
	HEMBB1000407	19.286	8. 572	18.593	3. 281	2.599	13. 454	2.473	
	HEMBB 1000420	95.847	66.573	138.307	54. 950	39.330	56.220	37.608	43.081
	HEMBB1000430	274.820	161.981	153.601	40.874	406.081	489. 107	693, 805	115.638
	HEMBB1000434	350.936	139.481:	599.497	199. 198	125. 426	113.500	65.776	77.687
	HEMBB1000438	67.342	10.187	25.472	7.736	8, 148	27.875	7.217	5.701
	HEMBB1000441	84.086	98. 109	312.643	78.842	60.934	76,141	46.589	35.267
20			88. 156	54.883	26.628	31.157	24. 328	25.777	38.008
	HEMBB1000447	76.519				1.700	13.053	5. 731	8, 109
	HEMBB1000449	22.357	11.282	25, 245	11.267				15, 902
	HEMBB1000453	26.781	29.875	49.056	22.139	35.305	22.456	14.006	
	HEMB81000455	37.937	43.401	129.423	29. 222	40.584	24.577	21.227	20.356
	HEMBB1000472	146.390	61.195	235.753	80. 306.	44. 122	82.882	52. 783	87.457
05	HEMB81000480	138, 135	67.904	194.466	46. 367	41.944	60.409	34.897	40.785
25	HEM881000486	78.511	63.045	211.876	47.786	39.049	36.558	20. 396	17.632
	HEMBB1000487	21.510	22.091	29.116	10,718	21, 056	15.854	13.086	10, 892
	HEMBB1000490	232.419	148, 116	562.064	159.218	134.370	107.861	60.296	110.306
			107.169	349.100	81. 342	44. 330	51.147	33.633	59.342
	HEMBB1000491	149.070				6. 821	10.632	9. 805	5. 454
	HEM881000492	18.194	21.930	19.080	9.690				
30	HEMBB1000493	286.390	34.074	64.876	31.406	23.065	49.816	39.824	39. 921
00	HEMBB1000510	133.225	95.239	380.177	165.002	101.728	72.504	64, 646	83.048
	HEM881000516	137.574	35.610	61.963	35.305	10. 932	78.851	39. 905	19.224
	HEM8B1000518	8.388	3.267	26. 133	5.489	1.531	1.500	1.611	1.901
	HEMB81000523	153.793	88.071	329.880	82.474	43.568	69.756	32.830	51. 127
	HEMBB1000530	46, 151	13.390	40,950	8.319	32.799	6.126	10.689	8. 425
		57.808	35.831	46. 332	20.306	19.414	5. 489	13.314	22.747
35	HEMBB1000542			79.169	22.945	10.597	23. 147	37. 266	20. 568
	HEMBB1000550	39.123	26.036				98. 169	55. 377	100.995
	HEMBB1000554	192.214	105.635	349.184	148.874	90.632			40.726
	HEMBB1000556	100.759	22.180	68. 289	37.737	35.176	41.190	47. 163	
	HEMBB1000564	101.412	37.586	144.386	37.463	27.344	59. 939	31.447	9. 452
	HEMBB1000567	361.516	76.515	125.177	66.960	83.698	221.216	145. 840	54. 204
	HEMBB1000569	53.847	46,712	54.356	18.197	23.752	36.942	31.264	39.479
40	HEM8B1000573	99.088	89, 487	373.557	76.986	76.236	63.534	39.587	58.648
	HEMBB 1000575	74.071	67.726	295. 427	63.469	37, 530	42.388	33.544	46. 151
	HEMB81000579	27. 868	12.805	18. 934	6.889	3,743	24.452	24.367	23, 262
				65.882	26. 172	19.828	26.184	16, 826	33.888
	HEMBB 1000585	30.826	34. 244					28.693	50, 228
	HEM881000586	85, 397	75.643	187.543	99.762	48.456	35. 430		
	HEMBB 1000589	135. 404	58.619	243.853	51. 181	36.284	29.883	21.561	27.997
45	HEMBB 1000591	99.680	60.946	242.306	54.695	36.589	52.616	32. 332	33.066
	HEMBB 1000592	30.320	18.740	34.338	11.753	8.732	28. 305	13. 707	12.164
	HEMB81000593	148.639	68.816	255.892	61.084	46.829	61.565	49.545	66.588
	HEMBB1000595	27.140	21.001	29.869		9, 199	21.841	16.487	29.680
	HEMBB 1000598		31.891	85.011		13.772	21.958	13.576	26.747
		39.074					15.614	9. 246	9, 161
£0	HEMBB 1000611	14.828	6.552					46. 273	77. 986
50	HEMBB1000617	193. 986	137.945				84. 583		
	HEMBB1000623	65.566	26.480				40.974	28, 571	23. 219
	HEMBB 1000630	62.606	23.074				31.973	21. 492	13.779
	HEMBB1000631	61.311	41.283	27.586	23.498	24.433	35.043	48.566	22.826
	HEM8B1000632	58.747	55.433				33.497	21.899	
	HEMB81000636	127.885	47.562					42.360	
55	HEMBB1000637	817.391	628.017				443.855		
55	LICTURE 100001	1011.331	1050.011	110-3, 130	1524.003	702. 301	1	1	

Table 42

	Diction Lange 20	55, 058	47, 453	95.751	42.262	25.684	15.056	22.121	28.829
	HEMBB 10006 38			251.754	80. 865	42.458	81.296	37. 596	52.009
	HEMBB1000642	179. 188		113.037	18.985	11.038	14. 245	8. 276	18.743
5	HEMBB1000643	43.411		137.371	34.816	24. 496	9. 967	11.881	22.322
	HEMBB1000649	27. 852			78. 131	42.090	36. 343	22.852	31.597
	HEMBB1000652	84. 942		126. 562	57.166	32.733	57. 424	38. 897	44, 477
	HEMBB1000655	418.308	73. 377	56. 858	20.817	6. 796	13, 110	7. 987	4. 458
	HEMBB1000665	16. 253	13. 954	10.766		25. 844	15.049	12.549	11. 202
	HEMBB1000668	28. 587	13. 435	14.606	13.788	96. 244	75. 058	66. 812	88. 267
10	HEMBB1000671	239.020	122. 952	561. 221	119.970		4.811	2. 141	12.812
	HEMBB1000673	11.633	5. 779	14.629	14. 904	5.916 1.049	30. 246	7.774	7.063
	HEMBB1000679	16. 899	7. 357	23. 438	7.697	66.411	89. 722	49. 173	67.832
	HEMBB1000684	188.240	157. 754	430.254	128. 150	1. 158	3. 240	3. 421	1.785
	HEMBB1000692	4. 978	9. 265	11.569	5.085	25. 408	13. 898	31.488	20.706
	HEMBB1000693	63.119	40.561	59. 522	22.326	19. 928	11. 568	2.839	10.179
15	HEMBB1000705	15. 560	31.798	122. 757	36.451 8.621	11.683	41. 509	10.019	7.584
	HEMB81000706	22. 553	13.626	23.777	50. 833	51.093	50. 427	37. 955	51.357
	HEMB81000709	74.737	77. 864	245. 726	12. 298	6.349	9. 891	2. 142	14. 350
	HEMB81000714	23. 726	10.733	6.625	13.761	12.596	17, 372	8. 105	16.144
	HEMB81000725	24. 239	9. 575,	11.437 208.396	65. 157	43.881	37, 441	22.020	39.067
	HEM881000726	86. 971	25. 288	140. 931	23.005	27.775	18. 629	12.838	14. 902
20	HEMBB1000729	51.556	38. 955	166.616	42.588	21.380	43. 330	7. 181	21. 192
	HEM881000738	39.002	94. 942	454.741	136.454	54.340	39. 253	32.933	49, 141
	HEM881000749	115. 917 47. 835	25. 201	36.488	16. 952	21.036	31.919	14.990	12, 111
	HEMB81000763	30.598	45. 410	167.003	32.786	26.482	25.698	18. 186	24. 127
	HEMB81000774	27. 168	21.690	33.470	20.937	12.916	22. 598	8.092	17.606
	HEMB81000777	245. 286	57.131	58.743	31.851	40.345	119, 113	81.364	53.990
25	HEMB81000781	41.945	36, 620	34, 149	24. 543	23.561	16, 383	14.371	20.775
	HEMBB1000788	10.756	10. 808	5. 481	6.429	2.950	5, 995	4.522	4. 589
	HEMBB1000789	28, 490	9.620	26. 151	16.088	11.640	16. 477	7.916	7.672
	HEMBB1000790	74, 318	56.925	185. 959	63.749	33.523	24. 232	24.414	28. 423
	HEMBB1000794	18.080	17.254	38.875	24. 305	7.427	10.338	5. 445	9. 305
	HEM8B1000807	50.070	31.869	22.751	19.865	20.934	27. 002	18. 350	27. 280
30	HEM881000809	334. 541	42.976	42.300	26. 454	9.545	31.526	31.677	44, 152
	HEMBB1000810	189.365	50.676	163. 325	33.349	38.994	74. 400	45. 398	19.262
	HEMB81000821	40.710	9.304	21.006	6.841	5. 422	15. 981	10.835	5. 685
	HEMBB1000822	8.726	3.570	3.541	1,411	7.255	5. 519	1.285	1.525
	HEMB81000826	68. 485	40.348	201. 149	68.467	43.204	31.769	32.812	55. 367
	HEMBB1000827	50.671	34. 326	108.391	32.945	15.076	25.813	18.713	25. 457
35	HEMBB1000831	38.060	20.466	29.131	12.368	19.990	20. 562	25. 373	6.415
	HEMB81000835	59. 181	56. 345	127.358	58.150	44.350	35. 831	25. 687	35. 108
	HEMB81000840	117.639	63.375	340.802	61.186	48.924	38. 995	20.712	30. 576 29. 431
	HEMBB1000848	98. 938	53.024	210.423	42.569	28.984	47.603	29.642 1.364	1.086
	HEMBB 1000852	1,827	2.160	0.621	2.559	1.621	1.272	10.179	14, 915
	HEMBB1000857	16.897	16. 768	19.951	14. 921 8. 209	14. 482	12.749	92.823	10. 102
40	HEMBB1000858 HEMBB1000867	25.634 106.946	16.531	8. 162 264. 748	50.278	36.949	41. 202	26. 795	29.760
	HEMBB1000870	68.550	62. 423	192.351	52.406	39. 103	55.641	23.738	27.427
	HEMB81000876	21.813	12.044	24. 968	11.314	7.689	10.690	11, 143	26.241
	HEM881000881	30.089	16.478	28. 345	14. 926	18.419	17.763	18, 901	20.494
	HEMBB1000883	11.669	10.263	26. 185	6.975	2.780	8. 223	2.906	3.540
45	HEMBB1000887	42.638	32.274	66.780	22.979	31, 512	42.842	20.622	22. 566
45	HEMBB1000888	20. 318	8. 193	11.483	5.178	4.073	8,708	6.801	4. 342
	HEMB81000890	48.795	42.287	112.076	25.031	11. 171	23, 116	15.491	16. 447
	HEMBB1000893	38, 227	10.603	88. 306	24.535	14.440	12.863	9.734	17.727
	HEMBB1000900		8.709	17.013		10, 928	12.199	14.105	11, 108
	HEM881000905	63.589	43.501	37. 125		26.379	29.649	38.699	31.891
50	HEMB81000908		54.674			28.838	28. 194	15.897	26.230
	HEMBB1000910	72.960	51.795			36.594	37. 378	13.612	23. 263
	HEMB81000913		35. 219	96.448		12.371	26.067	14.715	19. 268
	HEMB81000915		222. 511	693.345			1548. 228	1159.943	223. 176
	HEM881000917	99. 538	64. 212				34. 989	22. 324	40.667
	HEM881000927		11.252				24. 548	17.769	17.391
55	HEMBB1000932	33. 128	33. 556	95. 029	29.041	17.945	21.758	22. 973	31.034

Table 43

	DEUDD (AAAA)	907 670 1	93.035	605. 052 12	89. 543	312.660	538, 431	353.155	291.706
	HEMBB1000933		17.243	46, 380	14. 205	25.527	13.908	8, 530	11.716
	HEMBB1000936	23. 212		52. 525	19. 304	30.111	35. 223	41.856	37. 146
5	HEMBB1000939	105.016	35.905	15.872	4.660	6.130	17.648	83, 246	9. 541
	HEMB81000941	6.540	27.555		21. 361	9. 565	34, 299	13. 482	13. 269
	HENBB1000947	36.384	18.020	19.810	11.407	6. 851	17, 302	10.023	8.877
	HEM8B1000954	16.970	17.878	78. 296	22. 443	5. 599	10. 268	10.673	12. 183
	HEMBB1000959	22. 223	21. 226	21. 189	8.579	7. 102	23.845	5. 510	9.891
	HEMBB1000973	11.584	10.364	69.027	23.084	27. 137	40. 162	56.997	30, 316
10	HEMBB1000975	99. 598	37.022		20.141	5.813	6. 152	13.771	4. 102
	HEM8B1000981	10. 199	12.524	7.574	4,776	6.642	2. 985	6.049	3.612
	HEM8B1000985	13.065	8.026		8.864	8, 560	28. 394	25,072	10.907
	HEMBB1000991	67.124	17.092	28.053 352.650	90. 350	64. 926	71.240	60.014	102.622
	HEMBB1000996	170. 256	127.636		15. 226	10.611	14. 541	7,698	9.642
	HEMBB1001000	48. 257	19.380	16.573	0.000	0.000	0.318	0.000	0.000
15	HEMB81001004	0.797	1.839	0. 439 16. 434	11. 194	6.400	12. 238	6.478	9. 235
	HEMB81001008	17. 533	13.975	28. 396	15.752	15. 302	17.720	15. 586	16.702
	HEMBB1001011	39.743	19.337 46.352	244, 715	50.619	33.004	55.708	30, 100	34. 469
	HEMBB1001014	121.726		243.352	67.763	53.522	50.406	30.247	49. 844
	HEMB81001020	86.065 66.546	68. 022 <u>1</u> 59. 010	205. 347	41.480	31.865	35. 052	21.045	39. 489
	HEMBB 001024	36. 265	27.027	76. 443	19.990	25.484	27.657	12.014	23.129
20	HEMBB1001026 HEMBB1001037	64. 392	37.810	120.090	20.652	22.459	27. 294	18.918	28.917
	HEMBB1001042	58. 936	20. 428	42.468	17.255	15.600	32.463	20.274	18.506
	HEMBB1001042	76, 790	22. 021	40.791	13. 932	17.825	47.853	26.672	30.056
	HEMBB1001048	76.665	39. 237	208.757	53.469	44.539	37.624	16.049	20.262
	HEMBB1001048	133.028	58. 176	140.515	48.390	34.614	42.111	29. 526	34. 858
	HEMB81001051	22.699	8.465	13.142	9. 942	10.085	9.946	5.881	8.790
25	HEMB81001056	40.040	16. 494	45,000	22.674	18.685	21.131	18.431	12.498
	HEMBB1001058	88.873	59.116	223.822	45. 122	34.696	29. 783	21.562	25. 222
	HEMB81001060	35. 486	18, 631	33.852	60.851	26.807	13. 499	12.993	19. 391
	HEMBB1001063	53.418	36.359	125. 166	33. 156	24.220	19.182	16.188	14.597
	HEMBB1001068	79.181	46.879	78.756	35.034	26.835	79.006	63.198	43.296
	HEMB81001082	66.296	58.491	173.393	49.675	25. 253	33.015	14. 189	22.904
30	HEMB81001095	64.435	31.409	20.825	17.116	14.939	41.581	21.497	13.792
	HEMB81001096	43.372	28. 562	94. 366	32.120	13.089	21.236	15.814	22.034
	HEM881001101	79.652	21, 131	40.775	18.757	35. 350	46.263	18.855	13.874
	HEMB81001102	51.740	27. 585	86.794	21.160	12.958	16.450	7. 235	8.605
	HEMB81001104	61.846	33. 489	28.997	14.789	10.623	20.859	15. 993 20. 779	10.658
ar.	HEM8B1001105	69.199	32.868	132.855	27. 292	32.605	49.984	87.696	95. 854
35	HEMBB1001112	161, 356	78. 361	73.588	64.617	86, 150	93.363	32.824	66. 952
	HEMB81001113	114.744	130.208	298.139	107.218	73.757 62.314	35. 251	34. 480	51.970
	HEMBB1001114	105. 358	95. 960	365.719	26.838	17.638	29. 948	23.803	34. 239
	HEMB81001115	67.274	16.815	13.190		4.937	2.694	2.729	18. 952
	HEMBB1001117	2.434	10.819	14.951	4. 152 15. 560	5. 202	13. 437	5.261	9.614
40	HEMB81001119	18. 198	17.501	58.077 266.365	81. 302	76.905	130. 782	58.863	61. 487
40	HEMB81001126	306. 301 39. 673	36.703	178.312	45.328	36, 363	38.712	14.400	26. 997
	HEMB81001133 HEMB81001137	53. 424	19. 209	46.849	14. 453	13.705	30.395	18.865	15.761
	HEMB81001142	105, 888	131.411	405.403	98.008	104.700	62.754	32.598	75. 485
	HEMB81001145	114. 864	106. 329	348. 161	78.364		54. 983	24.738	51.568
	HEMB81001151	149.618	23.632	66.607	14.582		68.060	46.084	19.806
45	HEMB81001153	92. 263	53.444	153.351	44.131	37.191	34. 991	21.708	32.599
43	HEMBB1001158	64. 416	30. 844	50.578	22.880		47.046	24. 553	39.658
	HEMB81001169	96. 424	70.158	253.814		TAL ALA	37.113	24. 102	38.757
	HEMB81001170	34. 989	7.730	32.617			11.418		5. 208
	HEMBB1001175	46, 512	27, 401	45. 252		15.416			36.021
	HEMB81001177	126.389	86. 212	396.633		48.470			
50	HEM881001182	70.825	30.508	45.077					26. 488
	HEM881001192	30.059	21.703	61.610		5.688			
	HEMB81001199	1.469	0.000	0.000					
	HEM881001200	2. 266	1.426	2.071					
	HEMBB1001208	111.969	37.738	122, 154					
	HEMB81001209		77.445						
55	HEM8B1001210	14.499	40. 527	32.902	6.231	10.125	16.413	17.251	28.930
				_					

Table 44

	HENBB1001215	219, 922	83.033	126. 326	63.007	71.733	115.441	61.961	72. 230
	HEMBB1001217	63,633	22.116	41.047	17.479	20.160	53.164	31.645	18.739
	HEMBB1001218	98, 226		142.266	53.412	29, 467	23.819	20.495	24.079
5	HEMB81001221	0. 524	1, 310	12.795	0.988	0.992	0.867	0.000	1.767
	HEMBB1001224	52, 109	37. 281	86.318	28.364	24.177	19.072	16. 478	20. 321
	HEMBB1001230	38, 785	17.158	30.714	15.256	12.698	31.469	27. 596	17, 436
	HEMB81001234	335, 966	64.817	131.669	43.601	69.385	167. 134	101.415	57.258
	HEMBB1001235	152,870	67. 952	84.726	40.262	26.665	52.686	38, 623	49.693
	HEMBB1001237	16, 971	23.623	33.663	30.744	21.161	18.495	18.264	25.643
10	HEM881001242	26.787	15.776	22.922	4.200	5. 187	11.277	10,621	7. 589
	HEMB81001244	280. 439	9. 589	9.743	8, 128	2.116	4. 366	2.735	2.871
	HEMB81001249	51.892	27.766	106.010	25. 983	19.890	21. 254	16.839	21.542
	HEMBB1001253	50.869	33.773	58.857	31.656	8.253	38.144	20. 639 22. 558	25. 942 8. 559
	HEM8B1001254	28. 109	8.716	61.080	12.779	6.376	18.461	1.717	1.611
15	HEM881001266	2.010	9.088	3.704	1.682	16. 420 45. 610	62.418	33. 457	63.350
	HEMBB1001267	131.334	93.697	391.730	88.886	15.244	12.530	8. 683	10.739
	HEMB81001271	31.480	28.408	63.773	19.821	7.531	21.762	15. 737	10. 592
	HEMB81001282	41. 166	11.440	25. 546 131. 870	63.454	15. 491	70.758	43. 360	52.931
	HEMBB1001287	195. 274	10.227	25. 481	9.789	5. 520	21.519	16, 538	9.861
	HEMB81001288	40. 232 84. 233	74,730	246, 417	61.615	31.689	36. 447	24. 521	38.077
20	HEMBB1001289 HEMBB1001290	57.742	13. 181	11. 174	33.921	23.320	24.860	82.615	15. 369
	HEMBB1001294	80.761	23.745	72.937	16.689	20.147	45. 268	37.686	22.951
	HEMBB1001299	58, 616	17.094	44. 424	13.532	14.650	31.325	32.822	12.329
	HEMBB1001302	87. 107	24.979	56. 357	23.389	20.784	37.921	28.849	21.981
	HEM8B1001304	12, 134	0.119	5. 246	19.403	1,810	3.978	2. 153	1.580
	HEMBB1001314	6.410	5.111	25.042	5.961	3. 244	7.037	2. 954	2. 258
25	HEMBB1001315	3.706	8.398	10.733	3.067	1.405	3.652	1.659	1.943
	HEMBB1001317	39, 137	34.918	87.084	32. 290	25. 473	21.551	14.009	18.118
	HEMB81001326	13, 902	5.726	7.704	2.886	2. 324	1.546	2.008 24.975	5.612 17.786
	HEMBB 1001331	34.871	17.866	37.859	11.626	6. 188 15. 964	18. 477	15, 941	5.614
	HEMBB1001335	22, 550	20.911	19.341	12.458 45.250	52. 185	20. 178	25, 750	29. 233
30	HEMBB 1001337	61.645	43.894	187.675	11.541	12.874	18. 490	12.601	13,466
50	HEMBB1001339	31, 209	25.030 8.322	15.710	5.412	6.749	16. 517	16.482	9.869
	HEMBB1001346	44. 149	21.512	38. 191	15.415	9.432	26. 936	17.706	15.965
	HEMBB1001348	66.624	40.319	173.356	39.887	28.835	31.783	20.641	26.670
	HEMBB1001350	103.603	17, 400	35. 832	13.555	13.837	54.503	34.694	19.925
	HEMB81001356	12.440	11.385	25. 095	8. 592	6.787	7.806	8.759	8.923
35	HEMB81001364	28. 525	14.483	31. 452	11.829	13.494	12.620	13.025	10.117
	HEM8B1001366	57.883	53,690	210.263	52.112	27. 208	41. 191	29.156	32.064
	HEMBB1001357	140.660	59.744	283. 101	54.260	46.338	67.368 12.537	7, 158	48. 485 14. 407
	HEMBB1001369	17. 341	20.708	71.044	14.855	7.629	41.591	29. 026	63. 358
	HEM881001380	50. 204	67.647	124. 463	14.113	43.730 18.710	9. 428	10. 202	13.801
	HEMBB1001381	19.588	19.545	34. 218 26. 926	11.606	19.030	10.038	7.367	14, 535
40	HEMBB1001384 HEMBB1001387	17.779	16.837	19. 148	9.955	8.901	15. 994	7.831	13.345
	HEMBB1001394	21.419	19.091	32.720	17.551	19, 172	11.590	12. 282	11.322
	HEMBB1001407	39, 158	17.718	75.721	24.299	17.481	17.410	20.342	15. 525
	HEMBB1001410	18.880	3.346	6.042	2.907	2.655	0.000	2.839	2.094
	HEMBB1001413	32.291	25.769	80.279	17.033	21.102	11.132	12.610	24. 207
45	HEMBB1001419	36, 323	42.415	185. 239	24.790	21.849	17.972	13.895	31.342
	HEMBB1001421	29. 464	57.495	109.370	12.065	15, 685	64. 181	165.647	23. 322
	HEMBB1001424	9, 663	7.148	10.294		6.773	7.183	5. 215	8. 524
	HEMBB1001426	36. 471	25.897	86.872		17.823	19.534	15. 347	23.782
	HEM881001429		47.669	19. 928		21.695	39. 456 86. 724	39. 474 48. 813	58.527
	HEMBB1001436	168. 445	86.814	350.902		54. 546		9.581	12.477
50	HEMBB1001443		11. 137	12. 445		16.707 23.391	14.531		22.377
	HEMBB 1001449		34.064	145.511		28.709		29.720	26.623
	HEM88 100 1454		40.766 28.808						28.079
	HEMB81001458 HEMB81001461	77, 938							27.974
	HEMBB 1001463								38.517
55	HEM8B1001464								2.964
55	112201001101	1							

Table 45

	DEUNDAGA (EE	21 340	22. 324	20, 480	15.496	3.611	15. 533	10.020	13.761
	HEMBB1001456	31, 340		9. 987	4. 175	4. 887	24. 039	4.114	4.470
	HEMBB1001482	12.741	4.057		17.492	9, 196	12. 958	8. 167	14.603
5	HEMBB1001500	26.823	21.417	65. 107	04. 682	36, 419	54, 346	38. 027	46. 591
	HEMB81001505				25.792	20. 204	23. 504	18.628	22.786
	HEMB81001521	55. 379		133. 188	131.308	116.694	179, 333	72.732	79.869
	HEMBB1001527	331.186			19. 457	7.662	20.875	31.031	23. 503
	HEMB81001530	24.722	25.693	57 090	34. 674	21.061	27.704	18,966	32.578
	HEMB81001531	43.913	51.679	130. 225		1.875	8, 172	300.808	7.501
10	HENBB1001532	6. 957	3.901	34. 322	7, 593		34.644	21.690	23.017
	HEMB81001535	71.654	59. 202	131.794	45.369	28. 936 16. 411	22.356	19.126	20.785
	HEMB81001536	73.109	48. 204	106.813	35. 175	21. 583	31. 273	8.692	29.500
	HEM881001537	40.809	54.756	140.043	43.830	26, 100	44, 300	19.679	22.657
	HEMB81001542	79.436	33. 152	94. 294	34.360		20.740	11.834	18.032
	HEMBB1001543	55.819	14.588	8.417	4.239	7.702	15. 047	10.631	8. 198
15	HEMBB1001547	10.745	8. 433	12.415	9. 202	26. 421	115. 789	76. 174	67. 211
	HEMB81001548	163.125	42.223	39.134	33.781	20.037		7078.074	11. 439
	HEMBB1001551	32. 248	10.176	8. 937	9.728	40. 731	30. 981	17. 189	40.721
	HEMBB1001555	52.998	58.959	166.842	57.865	18. 852	28. 472	27.682	23. 295
	HEMBB1001562	67.088	35. 544!	83.929	24, 475 304, 052	124.857	300.720	202.502	439. 361
	HEMBB1001564	139.467	320. 422	580. 390	39.891	29. 530	30, 029	17.527	28.501
20	HEMBB1001565	56.749	43.545	123.727	28.883	16.462	19.020	8.403	16.605
	HEMB81001569	34. 482	26.904	100.487 65.598	41.979	32. 247	35. 238	25. 583	36.979
	HEMBB1001573	48.940	40.308 57.831	211.685	61.076	40.832	38. 446	18.915	42.636
	HENBB1001585	153.364	40.343	113.224	34. 425	18.386	24. 673	16.535	26.124
	HEMBB1001586	44.946	130.811	402.650	111, 293	69.831	80. 240	46.050	75.499
	HEMBB1001588	157.947	11.160	44. 464	13.949	6.811	11.538	4. 359	11.569
25	HEMBB1001595 HEMBB1001596	53.986	20.798	39.629	25, 473	20.578	32.621	23.309	36.564
	HEMB81001599	29.275	7. 352	13. 267	11.568	5. 279	15.756	10.260	5. 135
	HEMBB1001603	3.581	2.642	7.782	4. 279	3.051	0.341	1.424	3.160
	HEMBB1001606	6.897	7. 220	7. 226	7.657	3, 104	5.383	5.658	4.364
	HEMBB1001612	101.576	58. 128	240.469	58.770	36. 287	42.917	27.221	40.063
	HEMBB1001618	52.604	38.648	141.745	37.723	24.274	24. 922	17, 197	24. 223
30	HEMB81001619	59. 431	78.268	138.545	63.285	52. 275	37. 035	22. 185	38.081
	HEMBB1001623	33.128	8.489	11.122	6.318	8.326	16.007	3. 331	7.918
	HEMB81001625	10.068	16.076	8. 496	7.577	2. 293	8.389	1.716	4. 647
	HEMB81001630	7.144	5. 464	31. 186	8.383	3. 256	11, 196	3.053	5. 942
	HEM881001635	18.151	8.186	33, 138	13.501	9, 143	9.688	44.037	8.859
	HEMBB1001637	40.224	35.174	58.964	24.082	26.640	26.340	20.792	26.243
35	HEM8B1001641	21.655	10.768	33.553	9. 122	5.845	7.210	5.796	8.300
	HEM881001653	76.468	45.984	138.114	33.606	30.023	33.136	16.720	25. 949
	HEMBB1001665	3.000	0.352	5, 654	0.275	0.718	0.106	0.899	0. 407 15. 687
	HEM881001666	48.027	23.276	59.669	22.201	9, 196	20.512	10.659	8.738
	HEMB81001667	2. 570	7.909	3. 107	5.847	8.690	2.748	1, 999	3. 932
	HEMBB1001668	2.545	8.886	13. 392	8.498	18, 131	3. 355 5. 110	4.341	2. 139
40	HEMBB1001669	5.751	5. 364	10.395	3.219	4. 970	22.401	12, 909	20. 514
	HEMB81001670	17.795	10.903	34.891	20.715	21.640	40. 433	25.038	49.339
	HEMB81001673	69.924	44.194	58.806	53.036	9. 356	22.270	15.894	11.977
	HEMB81001675	58. 961	9, 166	21.648	6.718	11.101	24.642	13. 266	4. 383
	HEMB81001679	51.245	11, 218	30, 139	14.666	11.548	25. 422	15.072	13.683
4-	HEMBB1001684	27. 854		34.446	7, 134		1.316	3. 180	6. 172
45	HEMBB1001685	9. 626	8, 721	4.741	1. 162		1. 109	1,036	1, 119
	HEMBB1001695	2.706 116.774	4. 723 37, 756	115.693		34.790	69. 383	44, 901	43.576
	HEMBB1001703		52, 606	211.228			43. 432		54.662
	HEM881001704	67. 385	70. 475	227.746			53,010		56.789
	HEMBB1001706	122. 282		154. 286			50. 260		43.746
F0	HEMB81001707	111.416	69. 815 16. 260	60.454					8.073
50	HEMB81001717 HEMB81001731	14.112 29.550	36. 222	21.992			35.654		32.089
		75.818	39, 477	107.419					17.320
	HEMB81001734 HEMB81001735	63. 245	22.136	169.823				17. 292	27. 924
	HEM881001736	20.722	18.061	27.944					
	HEMB81001747	21, 158	15. 281	18.501					
55	HEMB81001749		90. 342						
	(								

Table 46

	Company Andrea	-AP 10P T	64 646 T	101 001	44.766	60.100	45.138	48. 988	44.990
	HEM881001753	85. 135	63.020	101.881		42.763		32.005	30.821
	HEMBB1001756	86.556	37.048	83. 531	33. 276		54. 273		
_	HEM881001757	1.981	3. 522	5.232	3. 590	1.394	7.486	3. 256	3.014
5	HEMB81001760	13.573	14, 554	27.053	7. 204	5. 280	8.129	5. 242	4.088
	HEMBB1001762	26.210	15.945	24.826	8. 467	6.461	26. 934	6.893	9.656
	HEMBB1001780	18, 738	33, 363	27.562	17. 311	13.893	4.277	14. 584	19. 429
	HEMBB1001785	3. 266	2, 954	7.974	3. 522	3.900	7.429	3.964	4.008
	HEMB81001788	77.710	51,716	232.298	72.096	40, 555	41.418	29. 586	33.423
		221.348	29. 215	45. 528	20. 500	22.918	33.927	36.095	25. 245
10	HEMBB1001793		9.015	10.442	4.015	2.532	8.773	2.904	6. 333
	HEMB81001797	4. 049			14, 183	17. 392	26.448	23.001	29.744
	HEMBB1001802	430. 563	24.213	34. 832			54, 459	15, 772	55. 255
	HEMBB1001812	91.804	71. 389	218.174	56. 457	56.645	289.852	148. 011	
	HEMB81001815	506.853	426.652	275.995	120.005	129, 468			122.368
	HEMBB1001816	90.696	55. 478	178. 334	52.637	25. 170	45. 331	35. 194	47.899
	HEMBB1001831	22.874	14.551	46.474	16.825	9.329	19. 975	9.745	18.634
15	HEMBB1001834	456.615	299.793	406.927	241. 146	284. 283	499, 103	267.485	306.611
	HEMBB1001836	138, 292	91.459	348.309	101.544	73.058	67.103	40. 539	76. 261
	HEMB81001839	9. 720	6,600	7.318	0.000	2.606	4.296	2.217	2. 738
	HEMBB1001841	345.524	134.230:	67.049	25. 938	60.560	21.530	21.177	18. 485
	HEMB81001844	61.041	25.820	34.819	14. 237	14.648	34. 333	20. 655	31. 102
	HEMBB1001847	126.241	111,341	239.722	147.873	65.849	86.164	47. 980	108. 378
20	HEMBB1001848	40. 802	39.856	24.837	12.646	9.727	18.893	18.093	17.754
	HEMB81001850	171.151	101.141	118.680	33. 622	64.050	118.364	50. 599	75.857
			77.853	231.163	65.024	41.660	123.173	103. 961	48. 695
	HENBB1001859	133.676		255.141	83. 601	85.833	53.693	30. 832	49, 888
	HEMB81001863	115. 353	92. 421	8.336	10.061	4. 673	8.415	6. 299	9.816
	HEMBB1001867	15. 427	15. 822			8.810	8. 133	10. 520	11. 923
05	HEMBB1001868	24. 470	17. 457	24.238	7.996		45. 547	29. 853	39.008
25	HEMBB1001869	82.894	76.711	234, 322	61.007	44.801		9. 542	5. 964
	HEMBB1001872	15. 921	7. 288	5. 998	10. 151	2.561	5.674		
	HEMBB1001874	36, 336	11.065	22.113	15. 221	9. 515	14, 138	6. 058	5. 891
	HEMBB1001875	7,615	19.234	13.755	26.314	11.646	3.662	5. 863	7. 228
	HEMBB1001880	107.638	82.806	115.014	59. 163	39.712	47.440	27. 454	37. 214
	HEMBB1001899	15.785	11.630	15.181	7. 571	2. 259	12.203	4. 190	3. 366
30	HEMB81001903	59.215	24.149	27.564	15. 205	8.601	28.805	15. 592	15.765
	HEMBB1001905	29. 932	24.402	20, 256	15.117	8.559	17.138	12.021	12.009
	HEMB81001906	15, 456	13.077	51.260	10.147	16. 547	10.906	7. 943	9. 129
	HEMBB1001908	35, 095	32.316	100.465	26.514	24.742	20.649	8.759	14. 223
	HEMB81001910	67.419	35. 922	139.126	58. 266	43, 100	26.178	19. 330	29.710
	HEMB81001911	50. 456	46, 682	196.311	58. 337	31.782	35. 278	19. 934	32.009
35	HEMB81001915	40.796	27.017	19.351	20.885	15.345	12.662	9.798	36.052
50	HEMBB1001921	95. 398	115, 190	314, 157	85.049	59.940	59.397	36.034	60. 585
	HEMB81001922	54. 587	37. 299	107.814	29.796	15.712	23.741	15. 662	16.568
	HEMBB1001925	35. 478	39, 156	106.631	23. 241	15.055	16.405	13. 936	15. 471
	HEMBB1001930	9. 272	7.467	11.545	7.045	3, 402	5.636	2. 969	5.808
	HEMB81001944	122. 259	83, 163	268. 572	86. 582	66.995	51. 236	27. 262	45. 542
	HEMBB1001945	55. 555	20,668	28.702	7.169	21.076	24. 208	18.042	10. 472
40	HEMBB1001947	47.254	12.987	21.887	16. 223	6, 133	25.673	16.697	13, 440
	HEMB81001950	99.345	31,711	42. 202	32.724	17.168	68.211	28, 763	30. 429
	HEMB81001952	67.117	40. 169	164.691	39. 168	16.287	31.103	11.276	24. 511
			47.572	147.635	34. 659	22.662	21.660	13.445	22. 280
	HEMBB1001953	56.049	20.350	106. 261	26, 369	15. 837	16.589	5, 199	12. 837
	HEMBB1001957	43.669			48.003	21. 477	24. 564	17. 194	36. 361
45	HEMB81001959	26.731	45. 573	72.402		52.786	46.598	20.834	29. 320
	HEMBB1001962	59.585	38, 413	125, 747	48. 471		70.066		
	HEMBB1001967	155.252	96.306	460.639		89.090		46. 606	68. 839
	HEMBB1001973	62.418	55.111	203.353	61.777	40.564	39, 531	24. 193	43. 482
	HEMB81001978	205.611	67.998	184.804		42.195	56.711	62.043	55. 171
	HEMBB1001983	115.219	97.908	189.950			62, 957	41. 995	65, 291
50	. HEMB81001987	23.094	30.009	63.743		10.970	10.414	5. 543	10.645
	HEMB81001988	26.549		71.399		11.631	11.873	6. 563	10.248
	HEMB81001990	61.049	28.808	125, 791	31.477		26.525	9.894	24. 366
	HEMB81001996	40.435	12.303	17.096	14, 159		18.573	11.696	13. 433
	HEMB81001997	91.453	62.313	247.838	64.724		29. 522	27. 492	42.942
	HEM881001999	28.583			34.520	11.455	23.048	14.798	25. 158
	HEMBB1002002	19.354					12.044	7. 088	14.724
55									

Table 47

	HEMBB1002005	127. 202	87, 407	314, 165	82.406	66.505	55. 577	40.792	δ4. 185
	HEMBB1002009	0.000	1.364	22.770	0.807	4. 369	1. 295	0.000	0.000
_	HEMBB1002013	28.258	13.676	16.813	10.399	10.765	17.046	7.782	9. 691
5	HEMBB1002015	105. 576	48. 524	66, 937	36.377	38, 220	74.637	28.221	34. 621
	HEMBB1002024	216.724	27, 841	16, 159	12.961	10.268	16.725	13.378	30. 580
	HEMB81002035	46.139	20.267	93.090	25.830	19. 155	14.290	9.089	10.861
	HEMB81002039	56.819	33.510	91.779	23.686	12.816	13.451	13.710	16.666
	HEMBB1002041	64.639	34. 426	51.061	22.611	27. 241	31.364	25. 209	28. 240
10	HEMBB1002042	108.989	70.262	244. 087	61.596	54. 097	58.195	45.407	53. 478
.0	HEMBB1002043	45.022	36.752	179.777	48.242	21.779	25.603	30.919	28. 446
	HEMBB1002044	13. 181	2.012	5. 797	1.053	1.982	1.313	3.432	2.045
	HEMBB1002045	289.530	197.322	441.790	143. 182	150. 349	206.083	108. 290	118. 515
	HEMBB1002049	35.193 37.095	24. 481	83.015 49.110	26.999	19.710 13.580	27. 535 16. 690	16.278 9.422	24. 921
	HEMBB1002050 HEMBB1002051	36.389	16.954 19.655	68. 218	12.868	8,800	22. 352	16. 403	14. 540
15	HEMBB1002068	75, 935	30, 174	53. 312	27. 588	23, 758	28. 553	40. 522	36.664
	HEMB81002069	213.038	176.212	471.114	127. 141	113. 252	145.813	82. 555	84. 929
	HEMBB1002075	42.631	31.316	161.071	28.782	21. 239	25. 996	13.087	18. 589
	HEMB81002079	16.958	10.592	15. 974	7.658	4.913	11.054	12.406	9. 170
	HEMBB1002080	43.775	32.579	72.576	24.001	9. 827	28.608	17.214	17.433
00	HEMBB1002082	26.775	8. 257	21. 193	4. 448	6. 280	19.090	464. 903	8. 346
20	HEMBB1002084	17, 127	6.840	43.925	4.043	9.757	26.316	9.627	5.512
	HEMBB1002088	90.318	38.977	65.816	40.755	47. 974	81.367	57. 452	75. 281
	HEMBB1002092	192.949	59.522	268. 965	49.978	47.797	60. \$95	48. 524	38. 080
	HEMBB1002094 HEMBB1002103	127.875 29.830	84.707 9.307	379.671 18.867	89.066 12.419	80.779 117.011	70.636 11.825	38. 807 10. 555	57.037
	HEMBB1002109	28, 380	23.579	104. 568	24. 307	17.018	17.089	11.301	6. 133
25	HEMB81002115	71.073	86.440	117. 523	95. 976	28. 307	85.908	60. 445	114. 378
	HEMBB1002120	16.393	10.090	4. 147	2.085	3.568	9.594	4.954	4. 539
	HEMBB1002121	12.050	2.757	6. 522	1.146	2.007	0.000	1.999	1.549
	HEMB81002134	784. 781	365. 377	605. 805	262.168	223. 204	719.592	534. 370	450. 949
	HEMBB1002136	109. 220	32.405	75.010	27.402	26. 278	36. 231	38. 283	23. 593
30	HEMB81002138 HEMB81002139	17.812 51.267	14.057	17, 210	7.413	9, 287	10.613	20.319	9.644
30	HEMB81002141	82.369	37.549 29.424	168.617	14.566	17.855 15.214	27.091 39.768	16.428 33.139	22. 856
	HEMBB1002142	70. 553	42.309	156. 252	36.636	14.797	26.769	15, 277	22.894
	HEM881002145	40.661	16.263	15.725	8. 229	13.984	21.757	14.873	15. 525
	HEM881002152	46.728	36.893	105.608	65. 422	40.064	25. 225	29.211	42.935
	HEMB81002162	40.153	34.008	96.274	29.709	19.847	47.860	22.055	40.550
35	HEM8B1002173	53, 191	41. 151	147.055	26.912	34. 538	16.431	19. 449	25. 327
	HEMBB1002189	73.400	88.057	211.287	73.810	54.029	46.682	45. 749	55. 885
	HEMB81002190	33. 242 69. 174	51.561 22.324	233.972 33.672	10.803	19.665 18.423	27.376	13, 129	61.389 16.109
	HEMBB1002217	50, 175	37.602	98.092	38.769	24. 723	33.043	18. 735	39. 436
	HEMBB1002218	596, 902	272.867	712.867	191.461	186. 314	373.711	195. 571	197. 556
40	HEMBB1002228	88. 583	45.763	205. 932	47.852	46.693	41.923	37.485	53.876
,,	HEMB81002232	56, 752	32.790	128. 643	36, 535	28. 593	32.710	31.447	41.940
	HEMBB1002245	31.084	9.332	17, 943	11.049	11.834	11.864	17.012	14. 199
	HEMBB1002247	151.502	27. 325	64. 167	10.018	25.829	62.501	35.734	21.698
	HEMBB1002249 HEMBB1002254	153. 327	94.814	380. 989	29. 328	65. 579	80.049	62.653	85. 673
	HEMBB1002255	43.885 8.633	36.756 2.293	118.582	8.771	19.323	2.385	12.693 3.358	22. 229 3. 589
45	HEM8B1002266	5. 303	5.716	8,530	6. 222	1.842	2.404	4, 411	2. 295
	HEMBB1002271	160.682	46.654	157. 828	58, 291	63.843	72.913	62.659	73.702
	HEMBB1002280	24. 597	13.246	76.763	13.976	7.742	9.196	9.200	16.479
	HEMBB 1002296	67.004	21.270	52. 536	34.388	49.938	53.045	123.030	41.218
	HEMBB1002300	94.815	28.682	50. 102	35.939	13.923	29.792	25.246	21.629
50	HEM881002302	51.059	31.157	28. 441	17.568	17.905	26.026	22.516	30.501
	HEMBB1002306	35. 213	49.812	33.017	23.300	15.072	17. 296	14.490	16. 293
	HEMBB1002316 HEMBB1002326	19.773 201.896	8.638 126.797	19.354	3.667	9. 274 89. 356	9. 974 85. 970	8.613	6.883
	HEMBB1002327	85.792	48.221	184. 126	47.724	32.764	29.959	54. 052 17. 415	98. 198 34. 542
	HEMBB1002329	69. 191	21.714	43.746	25.618	17.775	24.892	32.481	27. 906
£ £	HEMBB1002340	18.233	28.462	7.730	3.702	3.055	4. 522	2.914	5.745
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Table 48

	HEMBB1002342	74.746	83. 579	169, 482	40.919	23.495	26.453	33.215	66.420
	HEMBB1002358	149.857	132.962	286.214	85, 160	50.855	67.646	36.624	78. 432
						44.093		35. 955	51, 139
5	HEMB81002359	160.804	77.260	219.199	68.995		58.049		
•	HEMBB1002364	102.885	74. 409	188.270	50. 973	55. 276	45. 770	40.780	59. 739
	HEMB81002366	152.074	77.016	248. 465	68.268	81.100	64.637	39. 912	60. 303
	HEMB81002371	44, 433	12. 342	26.565	13. 307	36.600	10. 553	9, 238	5.351
	HEMBB1002381	134.427	77.953	207.310	57.210	48. 215	64.049	51.493	77. 629
	HEMB81002383	164, 205	52, 312	94.064	31.346	31.368	30. 947	43.038	47.640
	HEMBB1002387	196.859	164.904	235. 139	49.485	25. 102	93.004	52. 536	43.092
10	HEMBB1002409	82. 986	49.978	112.097	29. 207	15. 402	37.667	36.064	38. 132
	HEM881002413	123. 367	87, 690	361.106	87.505	57.485	48.097	23. 254	49. 302
	HEMBB1002415	87. 091	31.703	92. 595	31.804	23.352	27. 293	21.815	24.444
				15. 995	5.848	21.533	16. 980	18. 246	25.253
	HEMBB1002424	13. 162	19.511			36.928	41.171	26.823	47.957
	HEMB81002425	84.086	69.689	238.147	82.198			38.610	47.517
15	HEMBB1002427	143.727	26.894	50.430	25.865	40.707	52. 937		
13	HEMB81002442	163.853	121.153	501.168	129.909	73.231	81.033	47.108	287.238
	HEMBB1002447	107.214	80.007	214, 338	58.963	41.313	60. 452	49, 159	44.523
	HEM881002453	163.250	93. 442	384. 443	93.027	68.808	58. 565	45. 254	58.810
	HEMB81002457	116.756	104. 520	330.657	83.026	48.720	50. 971	38.415	57. 991
	HEMB81002458	18.721	11. 278	23. 232	9.587	7. 205	6.051	4.659	4. 343
	HEMBB1002463	229.657	146.001	663.683	193.622	138, 458	104, 827	52.827	110.558
20	HEMBB1002465	44. 210	23.316	33.631	20.895	17.932	26.471	19. 122	19.703
	HEMBB1002477	98.948	27.813	153.875	11.062	36.071	16.072	13.791	8.347
	HEMBB1002479	23.249	59.003	73.224	14.014	10.084	13. 246	1.980	8.949
	HEMBB1002489	78.748	24.690	71.038	31,400	39.869	43.673	44.800	75. 957
	HEMBB1002492	9.080	6. 989	26.130	3, 092	1.453	5.608	1.415	2.381
	HEMB81002495	95.752	104, 949	301.328	60.728	72.404	45. 161	24, 771	61.121
25	HEMBB1002502	17. 132	17.866	14.643	16.170	15. 224	14.056	4.504	23.313
	HEMBB1002509	0.913	2. 235	7.269	4.304	0.743	1, 283	1.504	6.154
	HEMBB1002510	0.732	0.000	0.000	1.858	0.926	0.000	0.000	0.000
	HEMB81002520	249.875	127.604	585. 470	169.423	138.712	90.360	100.598	112.828
	HEMBB1002522	24.741	27.480	12.342	14.142	17. 452	5, 861	8. 292	8. 541
		63.012		87.388	45. 392	29.555	37. 187	25.642	36, 089
	HEMBB1002527		61.066	83.329	21.750	21.479	53. 227	440. 333	38.710
30	HEMB81002530	72.655	45.682		9. 953	5. 539	16.743	.11, 880	8.115
	HEMB81002531	40. 398	18.832	10.308			46. 591	28.712	37. 112
	HEMBB1002534	78. 552	49.139	154.741	66.211	30.154	13.078	23. 458	15. 919
	HEMB81002536	27.609	22.843	52. 264	17. 646	8. 234		9. 427	
	HEMB81002544	24.012	6. 185	27.814	13.117	39.363	15. 921		14.017
	HEMBB1002545	108.234	31.929	243.949	50. 972	16.032	40.343	31.828	13.472
35	HEMBB1002550	31.850	11.452	10.668	11.228	11.049	10.100	14. 262	14.910
	HEMBB1002556	125. 621	89.607	311.607	79.974	50.209	57.837	53.696	54.119
	HEMB81002571	33.047	21.526	54. 457	14.847	25. 892	21.961	5. 482	18.608
	HEW881002579	75. 252	55. 132	229.479	48.891	31.521	43.266	24.667	31.554
	HEMB81002582	100.572	56.574	258. 453	63.093	45.740	39.580	26.474	45. 912
	HEMBB1002584	8. 325	7.514	13. 574	6.883	1.796	7.555	6.183	4. 955
40	HEMBB1002587_	57.430	44. 383	60.900	47.981	30.048	30.562	19.161	20.854
	HEM881002590	114. 241	78.587	179.926	65.737	28.629	43.657	33. 101	34. 032
	HEMB81002596	278.617	90.944	275.018	69.006	68.247	114, 505	88.149	59.750
	HEMBB1002600	17.618	16.003	23.907	4.699	9.726	10.133	7.945	8. 940
	HEMBB1002601	67.910	48.188	183. 948	45.346	38.021	37.423	21.860	33.698
	HEMBB1002603	69.793	43. 222	141.343	36.733.	28.849	35. 264	22.033	29. 436
45	HEMBB1002607	64.941	36.284	134.598	39.424	22.220	31.501	15. 575	31.024
45	HEMB81002610	22.852	9. 200	51.294	16.832	6.664	12.856	6.433	6.515
	HEMB81002613	85.026	60.872	161.891	47.532	36.559	44.841	24.569	31.062
	HEMBB1002614	65.074	30.721	39.687	10.970	15.910	13.297	10.461	5. 438
	HEMBB1002615	230.370	55. 581	35. 517	11.758	7. 258	46.064	22.857	86.789
	HEMB81002617	69.016	67.288	254.296	42.530	30.217	36.395	21. 284	37.688
	HEMB81002623	92.506	78124	204.116	60.739	20.110	48.078	32. 253	43. 355
50	HEMBB1002624	77.755	27.026	163.976	33, 209	25. 309	20. 104	21.741	24. 486
	HEMBB1002631	10. 297	18.892	12.879	14.916	7.219	5.864	6. 990	11. 537
			68, 172	141, 149	41.853	40.290	23.649	21.781	44. 425
	HEMBB1002635	88.049				9. 200	38.890	35.668	29. 597
	HEMBB1002644	98. 956	65. 380	26.659		32.125		315.048	39. 477
	HEMB81002554	127.571	78.659	51.653			137.732		
55	HEMBB1002661	106.501	46. 651	47, 116	19.470	20.684	30.561	24. 281	118.028

Table 49

. 45

		100 700 1	42.600	100.008	36.841	24, 382	35.028	41,975	18.150
	HEMBB1002663	100.783		395.057	51. 235	74, 731	89.130	40.814	55. 262
	HEMBB1002664	179.828	131.008		4. 981	2.314	4. 033	3, 301	1.422
•	HEM8B1002677	2.206	3.466	5. 138	55. 886	44. 381	26. 944	27.017	42.278
	HEMBB1002683	118.247	69. 327	247. 117	17.772	9. 039	5. 460	8. 120	14. 377
	HEMBB1002684	40. 291	21.056	46. 317		3. 146	12.807	18.055	9.131
	HEMBB1002686	30.893	12.882	26.031	19.059	19. 960	25. 893	38, 755	15. 268
	HEMBB1002692	48. 969	24. 335	52.440	29.779			73. 429	19.005
	HEMBB1002693	129.760	76.886	322.740	70.620	62.314	7.613	10. 266	29. 797
	HEMBB1002597	41.673	38. 793	25. 105	8. 999	2.058	109.419	79, 393	99. 532
	HEMBB1002699	223.756	165.884		116. 529	77. 378		13. 253	
	HEMBB1002702	13, 506	15. 782	24. 367	3. 561	6. 434	15.699	18. 202	30, 589
	HEM881002705	29. 934	20.276	15.478	21.230	7.599	11.487		
	HEMBB1002712	29. 588	10.805	47.572	15.673	13. 434	15.691	7.559	16.536
	IMR321000028	77.081	39. 937	40.934	18.725	8. 281	41. 195	27. 733	21.472
	IMR321000031	50.644	21.357	34.754	22.184	15. 786	31.242	22.705	14. 148
	IMR321000034	76.518	63.230	37.290	51.243	23.808	43.858	26.605	67.455
	IMR321000039	66.895	68.027	83.136	36, 653	27. 339	62. 232	57.760	88.100
	IMR321000044	1.614	0.000	0,000	0.000	0.000	0.000	1.970	0.000
	IMR321000063	131.633	84. 822	δ6. 499	84. 753	43. 262	73.363	69.831	80.878
	IMR321000085	157.704	34. 180	42.747	11.752	50.766	66.106	54, 160	47.424
	IMR321000089	52.645	22.980	31.408	17, 365	13.731	36.296	27.222	10.181
	IMR321000091	39.993	32.664	43.895	41.311	25. 143	35.002	20. 444	63.906
	LIVER1000004	45,674	30, 112	69.445	16.874	11.073	28.505	106.044	24.660
	LIVER1000008	23.703	14.444	22.304	9. 381	15.657	274.778	344. 333	11. 282
	LIVER1000011	107.957	31.187	106.032	30.434	41.030	41.256	348.474	63.939
	LIVER1000022	402.839	177.843	270.232	82.143	125. 292	205. 780	141.934	124. 260
	LIVER1000025	61.584	42.776	172.307	36,300	26.856	33.045	34.820	42.189
	LIVER1000030	62.987	24.034	69.275	29.784	17. 581	22. 393	51, 178	22.556
	LIVER1000045	27.941	4.859	27.468	7.384	9.755	14.426	20.651	24.802
	LIVERIDOCO46	180. 297	117.998	24.240	23.527	16. 373	7.466	27.795	66,724
	LIVER1000072	24.097	35.964	6.976	11.158	7.657	8.260	16.555	4.898
	LIVER1000077	90.518	39.165	17.306	13. 193	25. 835	52. 139	348.056	37.506
	LIVER1000080	17.084	4.918	5. 980	9, 600	2. 294	5. 176	6.495	4. 479
	LIVER1000086	82.711	55.169	150.708	18. 858	19. 278	176.018	481.085	27.747
	LIVER1000092	61.883	36.836	116.592	27.330	16.805	25. 266	35.863	24.160
	LIVER1000095	54. 562	13.959	104.146	23.878	13.158	200. 163	137.395	5. 508
	LIVER1000097	138. 286	11.401	12.265	8, 127	9.389	9.669	32.751	7.159
	LIVER1000098 -	58. 055	39. 291	47.410	18. 991	19.124	20.338	142.508	19. 104
	LIVER1000100	81.693	64.546	94.504	29. 185	18.588	42. 254	23.727	58.633
	LIVER1000101	52. 507	16.303	57.500	10.286	8.662	17.642	6. 129	27.273
	LIVER1000106	46.259	32. 121	32.438	11.568	9.377	13.216	102. 126	16.904
	LIVER1000108	26. 277	50.565	62.172	25. 422	16.619	17.243	38.369	18. 508
	LIVER1000115	23.571	18.673	71.357	14.244	11.023	17.910	427.626	11,136
	LIVER1000120	100.902	21.640	35. 183	16.565	26.236	39.037	87.151	16.249
	LIVER1000138	69.624	27. 584	56.479	22.794	25.076	42.015	35, 937	23.833
	LIVER1000146	107.757	63.296	209.735	54. 534	42. 231	45. 210	254. 168	42.466
	LIVER1000148	141.467	42.327	108.510	37.031	31.920	62.584	125.466	65.728
	LIVER1000157	97. 282	37.198	50.979	49. 952	35. 021	43. 954	52.527	43.221
	LIVER1000161	100.902	24.883	57.647	28. 329	31.562	42.781	89.198	30.740
	LIVER1000167	97.214	29, 093	41.460	25.700	26.316	112.706	332.789	30.702
	LIVER1000174	53.927	23.440	26.353	13.595	12.625	36.580	71.460	10.512
	LIVER1000185	49.746	20. 428	31.630	13. 964	13. 391	16.773	16.676	14.878
	LIVER1000187	38. 332	8.211	15, 200	4. 654	8.084	9.846	567.808	8.320
	LIVER1000190	93.672	29, 635		15, 812				11.496
	LIVER1000192	141.875			32.936	41.210	79.500	128.608	47.907
	MAMMA 1 000009	99.036	77.266		72. 924	40.612	44.930	25.218	35.909
	MAMMA1000015	40. 458			13.017	12. 921	18.315	13.014	8.185
	MAMMA1000019	62.999			52.037	36.450	42.958	38.148	30.172
	MAMMA1000020	58.696			40.615	38.572	34.176	18.169	20.807
	MAMMA1000024	15.610			7.263	3.468	11.662	37.960	9.224
	MAMMA1000025	53.706	37.358		37.766	29. 177		18.530	
	NAMA1000043	170. 220	108,774	290.077	126.472	100.059		70.843	76.243
	MAMMA1000045	83.118			10, 125	5.779		7.895	
	MAMMA1000046	117.084	44. 858	285. 890	86.458	43.862	36.388	23. 428	22.376

Table 50

IAMMA1000055	65, 118	40.884	57.307	29.859	27. 445	33, 405	22.066	23. 563
IAMMA1000057	170.331	108.479	421.160	100.366	84.331	77.475	42.047	55.847
			153.319	49. 223	28. 927	42.539	25. 636	52.458
MMMA1000060	79.698	50.265						
MMA1000069	118.921	35.010	182.272	48.764	43.720	61.342	45.357	33.115
AMMA1000084	128.354	92.819	277.404	87. 542	63.176	65.262	34. 266	45.092
AMA 1000085	40. 199	20.019	40, 608	21.956	13.181	18.822	36.347	26. 209
					16. 101	26.961	15.531	22.390
MAMMA 1000092	77. 338	37. 915	167.474	43. 988				
AMMA 1000096	55. 344	38.495	38.888	25.605	11.893	44. 990	24.784	25. 160
MANA 1000097	62.546	54.694	52. 522	52.269	24.807	65,730	25. 787	23.298
				31.689	19, 430	26.892	22. 353	15.842
MAMMA 1000102	67.585	32.797	91.551					
MAMMA1000103	63.752	26.301	89. 530	30, 004	12.188	31.709	11, 461	14.718
MAMMA 1000106	37.916	23. 228	90, 795	22.075	14. 445	24.686	15.649	17.569
	58, 533	24. 502	43, 190	22.445	16.140	27.418	15. 487	13.269
MAMMA 1000117					24.039	42, 731	38, 062	43. 242
MAMMA 1000118	104.168	58. 433	63.822	8.833				
MAMMA 1000129	170.665	72.256	98.813	45. 970	22. 181	58.739	50. 197	14.587
MAMMA 1000 133	52.435	25.090	33, 061	20.713	14. 310	34.686	18.642	14. 101
			246.344	90.530	127.758	76, 596	45, 325	60.360
MAMMA1000134	106.522	79.090						
MAMMA1000139	78.566	47. 362	99.179	34. 535	22.772	37.601	28.841	28.280
MAMMA1000141	30, 121	20. 528;	28. 150	13.910	5, 510	14.314	12.120	15.748
		8. 669	41.797	8.690	9.949	10.059	4.040	8. 280
MAMMA 1000143	16.647							
MAMMA 1000150	128. 128	259, 413	21.844	28.777	86.623	42.827	51.840	42.986
MAMMA1000155	205.031	88.642	291.247	110.884	80.817	97.755	63.045	78. 585
MAMMA1000163	43. 643	36.898	57. 239	22.848	21.852	41.672	11.036	10.618
		46. 928	265.746	98. 189	60.007	66.037	34, 872	50.109
MAMMA1000171	141.225							
MAMMA1000173	103.027	21.955	68.080	33.572	25.668	45. 271	40.340	52. 609
MAMMA1000175	19.316	8, 683	7, 960	4. 550	3.535	7, 894	5, 974	4.015
MAMMA1000183	57.490	35.830	148.702	42.892	23.250	23.680	21.050	46.992
					22.392			
MAMMA1000191	88.722	31.449	40.834	26.064		26.766	36, 253	27.729
MAMMA1000192	53.467	25, 096	30.205	28.380	21.976	101.288	128.339	44.025
MAMMA 1000193	83, 936	36.823	36.836	29.409	18, 905	35, [3]	35, 059	36.667
			347. 292	70.840	49. 278	62. 924	38.858	66.720
MAMMA 1000 198	132, 127	93, 550						
MAMMA 1000204	64. 455	59.079	71.789	26.771	29.275	55, 156	62.132	49, 295
MAMMA 1000207	45, 771	62.052	52, 332	19.986	16.418	37.618	225. 196	18. 506
MAMMA 1000214	100. 292	62.311	289, 223	62, 541	32.825	57.748	32.755	39.770
				13.919	12.649	42.421	29.143	20.494
MAMMA 1000220	91.389	23.816	43.034					
MAMMA1000221	39. 338	35.655	11. 931	39.315	9.426	18.802	27.741	17. 121
MAMMA 1000226	65.096	20.174	11, 901	11.838	17.236	23. 487	43:016	24.801
MAMMA1000227	94.333	64. 156	183. 365		58.478	66.811	43, 961	53. 250
						56.380	71.726	37.727
MAMMA 1000230	116.378	47.908	97.869	47. 218	38. 196			
MAMMA 1000241	53.737	85. 177	107.748	60.815	31.230	51.839	36.525	22.770
MAMMA 1000245	107.413	148. 468	205. 437	144.478	51.682	86.017	93.183	198.398
MANNA1000248	205.478	88. 411	342.827	76.468	51.702	110.723	70.650	60.978
					42. 597	57.904	34. 572	51.015
MAMMA 1000251	115. 401	47.888	209.360	39.959				
MAMMA 1000254	43.161	20. 910	114.081	20. 548	9.699	9. 885	5.346	32.024
MAMMA 1000257	142.781	70.118	332.822	104. 425	84. 387	124.673	78.270	115.103
MAMMA1000262	18. 952	34, 301	19.786	32.516	14.840	15, 513	23.805	35, 519
MAMMA 1000264		20.630	124.043	44.847	29.466	21.390	22.616	37, 039
	59.532							
MAMMA 1000266	55. 476	28.959	122.654		27.018	24. 021	20.212	38, 284
MANIMA 1000270	142.968	64. 234	270, 948	75.022	64.760	68, 130	64,006	73, 994
	53.605	9.611	35. 682		16.139	24. 236	26.722	26.433
MANNA 1000271			98. 448	19.751	12.725	33.047	23.839	33.012
MAMMA1000271		16 125		1 13.131	1 14.143	1 33.07/		
MAMMA1000277	56.407	16.435		A 74A		70 475	1 25 20.4	
MAMMA1000277 MAMMA1000278	56.407 40.286	13. 365	19.395		12.609	20. 423	25, 204	
MAMMA1000277	56.407	13.365 36.984		46.809	12.609 34.441	42.500	26. 143	48.597
MAMMA 1 00 0 2 7 7 MAMMA 1 00 0 2 7 8 MAMMA 1 00 0 2 7 9	56.407 40.286 68.661	13.365 36.984	19.395 173.379	46.809	12.609 34.441	42.500	26. 143	48.597
MAMMA 1 00 D 2 7 7 MAMMA 1 00 D 2 7 8 MAMMA 1 00 D 2 7 9 MAMMA 1 00 D 2 8 3	56.407 40.286 68.661 55.199	13.365 36.984 27.095	19.395 173.379 46.168	46.809 22.395	12.609 34.441 15.870	42.500 21.308	26. 143 16. 298	48. 597 18. 504
MAMMA100D275 MAMMA100D275 MAMMA100D279 MAMMA100D283 MAMMA100D284	56.407 40.286 68.661 55.199 76.726	13.365 36.984 27.095 67.676	19.395 173.379 46.168 42.784	46.809 22.395 39.851	12.609 34.441 15.870 34.586	42.500 21.308 47.651	26. 143 16. 298 39. 169	48. 597 18. 504 48. 342
MAMMA 1 00 D 2 7 7 MAMMA 1 00 D 2 7 8 MAMMA 1 00 D 2 7 9 MAMMA 1 00 D 2 8 3	56.407 40.286 68.661 55.199	13.365 36.984 27.095	19.395 173.379 46.168	46.809 22.395 39.851	12.609 34.441 15.870 34.586 31.007	42.500 21.308 47.651 27.370	26. 143 16. 298 39. 169 29. 006	48. 597 18. 504 48. 342 35. 599
MAMMA1000277 MAMMA1000278 MAMMA1000279 MAMMA1000283 MAMMA1000284 MAMMA1000287	56.407 40.286 68.661 55.199 76.726 73.583	13. 365 36. 984 27. 095 67. 676 58. 726	19.395 173.379 46.168 42.784 142.953	46.809 22.395 39.851 39.301	12.609 34.441 15.870 34.586	42.500 21.308 47.651	26. 143 16. 298 39. 169	48. 597 18. 504 48. 342 35. 599
MAMMA1000277  MAMMA1000278  MAMMA1000279  MAMMA1000283  MAMMA1000284  MAMMA1000287  MAMMA1000287	56.407 40.286 68.661 55.199 76.726 73.583 457.450	13. 365 36. 984 27. 095 67. 676 58. 726 361. 106	19.395 173.379 46.168 42.784 142.953 313.407	46.809 22.395 39.851 39.301 116.696	12.609 34.441 15.870 34.586 31.007	42.500 21.308 47.651 27.370 343.951	26. 143 16. 298 39. 169 29. 006 155. 948	48. 597 18. 504 48. 342 35. 599 100. 375
MANMA1000277 MANMA1000278 MANMA1000279 MANMA1000283 MANMA1000284 MANMA1000287 MANMA1000294 MANMA1000298	56.407 40.286 68.661 55.199 76.726 73.583 457.450 31.731	13.365 36.984 27.095 67.676 58.726 361.106 25.511	19.395 173.379 46.168 42.784 142.953 313.407 41.413	46.809 22.395 39.851 39.301 116.696 16.220	12.609 34.441 15.870 34.586 31.007 112.848 16.320	42.500 21.308 47.651 27.370 343.951 14.676	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043	48. 597 18. 504 48. 342 35. 599 100. 375 20. 205
MAMMA1000277  MAMMA1000278  MAMMA1000283  MAMMA1000284  MAMMA1000284  MAMMA1000284  MAMMA1000289  MAMMA1000294  MAMMA1000298	56.407 40.286 68.661 55.199 76.726 73.583 457.450 31.731 109.379	13. 365 36. 984 27. 095 67. 676 58. 726 361. 106 25. 511 58. 532	19.395 173.379 46.168 42.784 142.953 313.407 41.413 280.880	46.809 22.395 39.851 39.301 116.696 16.220 69.156	12.609 34.441 15.870 34.586 31.007 112.848 16.320 44.790	42.500 21.308 47.651 27.370 343.951 14.676 36.788	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043 28. 220	48. 597 18. 504 48. 342 35. 599 100. 375 20. 205 40. 861
MANMA1000277 MANMA1000278 MANMA1000279 MANMA1000283 MANMA1000284 MANMA1000287 MANMA1000294 MANMA1000298	56.407 40.286 68.661 55.199 76.726 73.583 457.450 31.731 109.379	13.365 36.984 27.095 67.676 58.726 361.106 25.511	19.395 173.379 46.168 42.784 142.953 313.407 41.413	46.809 22.395 39.851 39.301 116.696 16.220 69.156	12.609 34.441 15.870 34.586 31.007 112.848 16.320	42.500 21.308 47.651 27.370 343.951 14.676	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043	48. 597 18. 504 48. 342 35. 599 100. 375 20. 205 40. 861
MAMMA1000277  MAMMA1000278  MAMMA1000283  MAMMA1000284  MAMMA1000284  MAMMA1000289  MAMMA1000298  MAMMA1000298  MAMMA1000303	56.407 40.286 68.661 55.199 76.726 73.583 457.450 31.731 109.379 67.505	13. 365 36. 984 27. 095 67. 676 58. 726 361. 106 25. 511 58. 532 14. 147	19. 395 173. 379 46. 168 42. 784 142. 953 313. 407 41. 413 280. 880 18. 804	46.809 22.395 39.851 39.301 116.696 16.220 69.156 11.073	12.609 34.441 15.870 34.586 31.007 112.848 16.320 44.790 33.859	42.500 21.308 47.651 27.370 343.951 14.676 36.788 26.599	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043 28. 220 30. 177	48. 597 18. 504 48. 342 35. 599 100. 375 20. 205 40. 861 30. 810
MANMA1000277 MAMMA1000278 MAMMA1000279 MAMMA1000283 MAMMA1000284 MAMMA1000287 MAMMA1000298 MAMMA1000298 MAMMA1000303 MAMMA1000303 MAMMA1000305	56.407 40.286 68.661 55.199 76.726 73.583 457.450 31.731 109.379 67.505 32.363	13. 365 36. 984 27. 095 67. 676 58. 726 361. 106 25. 511 58. 532 14. 147 19. 693	19. 395 173. 379 46. 168 42. 784 142. 953 313. 407 41. 413 280. 880 18. 804 108. 733	46. 809 22. 395 39. 851 39. 301 116. 696 16. 220 69. 156 11. 073 15. 375	12.609 34.441 15.870 34.586 31.007 112.848 16.320 44.790 33.859 12.695	42.500 21.308 47.651 27.370 343.951 14.676 36.788 26.599 14.455	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043 28. 220 30. 177 13. 353	48. 597 18. 504 48. 342 35. 599 100. 375 20. 205 40. 861 30. 810 15. 189
MANMA 1000277  MANMA 1000278  MANMA 1000279  MANMA 1000284  MANMA 1000284  MANMA 1000284  MANMA 1000294  MANMA 1000298  MANMA 1000302  MANMA 1000303  MANMA 1000305  MANMA 1000305	56. 407 40. 286 68. 661 55. 199 76. 726 73. 583 457. 450 31. 731 109. 379 67. 505 32. 363 279. 600	13. 365 36. 984 27. 095 67. 676 58. 726 361. 106 25. 511 58. 532 14. 147 19. 693 75. 098	19. 395 173. 379 46. 168 42. 784 142. 953 313. 407 41. 413 280. 880 18. 804 108. 733 397. 421	46. 809 22. 395 39. 851 39. 301 116. 696 16. 220 69. 156 11. 073 15. 375 75. 020	12.609 34.441 15.870 34.586 31.007 112.848 16.320 44.790 33.859 12.695 45.244	42.500 21.308 47.651 27.370 343.951 14.676 36.788 26.599 14.455 68.757	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043 28. 220 30. 177 13. 353 131. 117	18. 504 48. 342 35. 599 100. 375 20. 205 40. 861 30. 810 15. 189 116. 800
MANMA1000277 MAMMA1000278 MAMMA1000279 MAMMA1000283 MAMMA1000284 MAMMA1000287 MAMMA1000298 MAMMA1000298 MAMMA1000303 MAMMA1000303 MAMMA1000305	56.407 40.286 68.661 55.199 76.726 73.583 457.450 31.731 109.379 67.505 32.363	13. 365 36. 984 27. 095 67. 676 58. 726 361. 106 25. 511 58. 532 14. 147 19. 693	19. 395 173. 379 46. 168 42. 784 142. 953 313. 407 41. 413 280. 880 18. 804 108. 733	46. 809 22. 395 39. 851 39. 301 116. 696 16. 220 69. 156 11. 073 15. 375 75. 020	12.609 34.441 15.870 34.586 31.007 112.848 16.320 44.790 33.859 12.695	42.500 21.308 47.651 27.370 343.951 14.676 36.788 26.599 14.455	26. 143 16. 298 39. 169 29. 006 155. 948 22. 043 28. 220 30. 177 13. 353	48. 597 18. 504 48. 342 35. 599 100. 375 20. 205

Table 51

						60 666	T		
	MAMMA1000313	79.577	69.550	54.317	10.741	60.526	42.964	18. 206	37.303
	MAMMA1000331	80.910	48.868	139.047	33.811	22.564	·15. 207	18.580	21.385
-	MAMMA1000335	54.800	22. 399	33.190	18.244	16.273	30.688	26.611	30.790
5	MAMMA1000339	69.222	40.948	83.679	13.158	20.941	22.134	20. 026	10.739
					32.922	28.610	23.069	18.858	23.519
	MAMMA1000340	57. 498	34. 708	164.958					
	MAMMA1000348	78.099	102. 955	374.737	55.033	32.546	66.256	22. 303	23. 575
	MAMMA1000356	152. 238	116.086	454.516	67.232	34. 525	47.884	22.865	61.267
	MAMMA1000358	34. 367	56. 332	15, 362	15.091	16.743	17.405	19.645	7.358
	MAMMA 1000360	71.104	74. 351	246.244	43.414	24.093	24.945	14. 842	14.739
10								37.987	42.992
_	MAMMA 1 0 0 0 3 6 1	101.653	93. 468	230.215	73.577	45.022	37.236		
	MAMMA1000363	71.108	19. 232	39.013	13.717	23.713	30.739	27.813	32. 485
	MAMMA1000370	171.867	108.830	110.466	80.949	52.076	79.266	57.877	247.810
	MAMMA1000371	100.543	32. 223	80.873	48.039	49.442	91.739	57.647	46.599
	MAMMA1000372	206.850	114. 326	609,068	130.138	79.980	80.890	54.857	97.509
				238.462	40.061	34.528	31.361	22.458	45.681
15	MANMA1000385	72.074	60.911						
15	MAMMA1000388	118.855	69.094	105. 789	42.626	50.059	55.389	37.396	37.825
	MAMMA1000395	97.031	44. 493	34. 493	20. 201	19,036	27.695	24.269	17.433
	MAMMA1000402	126.085	107. 637	256. 584	68.415	45.669	61.486	30.340	30.943
	MAMMA1000403	87.558	63.749 €	208. 574	64.857	45.578	44.799	22.710	42.239
	MAJMA1000410	43.073	43. 539	94. 207	39.613	19.880	22.573	16. 272	21.003
	MAMMA1000413	30.829	13. 370	70.418	17, 102	13.392	15.291	11.599	15.353
20	MAMMA1000414	125.550	111.622	81.672	15.722	51.528	14.549	28. 214	13.858
			103. 793	427.214	107.383	105.899	121, 441	55.040	84.667
	MAMMA1000416	179.864							
	MAMMA1000421	131.712	73.475	307.780	70.841	55.037	49.498	34.519	46.482
	MAMMA1000422	12.614	14. 628	30, 167	16.100	11.675	22.441	18.843	54.831
							15.645		
	MAMMA1000423	34.100	22.150	69.677	18, 461	13.815		8. 500	8.869
	MAMMA1000424	9. 330	4.056	36.234	8.171	0.971	2.769	0.745	7.267
25	MAMMA1080429	575. 321	219. 503	317.414	158. 529	150.779	290. 300	196, 161	149.619
25									
	MAMMA 1000431	143.825	79.993	275.497	82, 499	52.496	63. 425	43. 337	66.733
	MAMMA1000432	65.212	17, 117	24, 472	28.083	17.360	33.881	27.547	29.615
	MAMMA1000437	89. 375	88.947	265. 572	50.025	69.885	45, 195	30.823	31.510
	MAMMA 1000444	120.017	124. 234	477.772	115.966	65.200	66.888	31.943	88.274
	MAMMA1000446	50. 201	66.027	41.406	8.991	18.971	29. 395	7.985	37.220
			41.427	180.761	40.414	25.983	35, 232	23, 109	27.942
30	MAMMA1000449	81.386							
	MAMMA1000457	47.862	13.862	15.095	11.981	7.566	21, 142	12.971	10.872
	MAMMA 1000458	34, 485	13.749	22.864	12.116	11.199	18.881	15. 924	10.046
						5.503	8. 258	7.138	
	MAMMA1000468	8. 235	7.843	6.029	5.004				1.618
	MAMMA1000472	250, 243	67.964	110.774	68.614	73.186	111.758	88.016	79.409
	MAMMA1000473	54. 174	16.506	40.489	16.002	17.450	26.506	17.741	13.900
35	MAMMA1000477	77. 316	50. 237	238.943	56.450	38.807	32.776	36.438	35. 332
	MAMMA1000478	201. 299	157.097	496.514	127.872	82.832	77.444	49. 296	86.763
	MAMMA1000483	107.340	74.564	252.463	60.824	31.055	44. 198	44. 157	87.449
	MAMMA1000490	14.473	14.068	16.023	12.496	8.202	15.654	11.091	12.344
	MAMMA1000496	32.756	10.554	20.693	10.676	19.830	19. 282	13. 204	13.410
	MAMMA 1000500	23.016	17.584	49.151	15.706	13.914	19.063	11.094	22. 904
40	MAMMA1000501	196.637	102.490	468.793	104.118	67.761	83.834	76.446	86.912
70	MAMMA1000503	7. 083	4.085	3.866	1,004	1.005	3.752	4.005	3. 248
	MAMMA1000506	201.452	116.279	151.434	56.847	78.502	149.780	99. 352	64.069
					39. 187				
	MAMMA1000510	70.898	18, 432	60.927		33.327	42.829	40.993	33. 127
	MAMMA1000515	43, 923	30.031	85.637	35. 744	18.805	21.837	19. 339	17. 922
	HAMMA1000516	74.742	48.811	148.307	43, 452	18.069	34.061	19, 122	26.985
0.=	MAMMA1000522	53. 273	23.845	132.197	22.861	14.594	24.776	12.095	27.578
45	MAMMA1000524	130.806	61.389	266.529	71.558	50.972	73.691	47.484	55. 510
	NAUNA1000528	38, 579	27.136			15.860	29, 316	19.300	24.797
	MAMMA1000534	32.603	20.088	33.950	10.973	7.185	10.580	7.972	10.160
	MAMMA1000541	165, 518	58.806	85. 648	63. 188	27.705	52.036	46.200	39.018
	MAMMA1000550	119.597		41. 184		5.859	48. 433	766.194	63.005
			203.059		24. 393				
	MAMMA1000556	31.963	15.056	15.588	8.634	11.294	15.698	21.467	16.597
50	NAMMA1000559	57.738	31, 181	242.155	29. 443	19.030	26.908	13.520	41.571
•									
	MAMMA1000565	118.770	30.318	289, 829	37.509	33.728	38.720	18.344	25.847
	MAMMA1000567	77.050	44.379	224. 645	48. 804	41, 102	56.039	36.496	63.529
	MAMMA1000576	271.038	180, 600	661.566	221.987	157.443	132. 385	93.679	129.843
	MAMMA1000582	54, 936	43.406	272.366	14. 342	18.896	29. 396	46. 333	40.210
	MAMMA1000583	90. 692	51.670	147.946	34. 905	17, 175	23. 177	19.077	40.824

461

Table 52

MANNA1000585 MANNA1000587 MANNA1000591 MANNA1000594 MANNA1000597 MANNA1000605 MANNA1000612 MANNA1000614	89.865 47.955 77.705 194.593 496.923	50.008 14.789 38.280	288. 673 58. 279	52. 259 12. 415	29. 243 6. 584	39. 188 14. 410	24.088 15.734	46.734 6.826
NAMMA 1000587 NAMMA 1000591 NAMMA 1000594 NAMMA 1000597 NAMMA 1000605 NAMMA 1000612	47.955 77.705 194.593	14. 789 38. 280	58.279	12.415	6 584	14 410	15 774	C 047
MAMMA 1000591 MAMMA 1000594 MAMMA 1000597 MAMMA 1000605 MAMMA 1000612	77.705 194.593	38, 280					13.134 (	0.020
MAMMA1000594 MAMMA1000597 MAMMA1000605 MAMMA1000612	194.593			28.019	20.094	28.578	24.299	19.949
MAMMA1000597 MAMMA1000605 MAMMA1000612			81.784				43.577	75.029
MAMMA1000605 MAMMA1000612	496, 923	94. 384	488.898	91.064	59. 244	55.681		
MANNA1000605 MANNA1000612		264. 906	751.636	196. 294	121.483	306.397	199.988	160.426
MAMMA1000612	324.584	183.667	990, 246	209. 555	135.844	158.096	97.598	149.183
		22.051	42, 999	14.074	19. 294	41,220	29. 460	15.713
MAMMA1000614 I	68.113			69. 022	127.808	309.892	249.344	194.110
	580.099	136.874	402.890					
MAMMA1000616	2.590	15.442	13.809	1.109	3.011	7.500	3. 036	3.188
MAMMA1000621	19. 258	12.723	14.307	13.200	5. 971	12.028	11.561	11.081
MANNA1000623	60.189	23. 285	25.913	12.057	10.648	23. 327	19.218	20.667
		249, 117	346.876	155. 944	192.671	373.924	300.473	274.263
MAMMA1000625	651.334					4.118	5. 584	9. 542
MAMMA1000635	4. 459	2.994	4.756	2.883	0.000			
MAMMA1000643	24.259	51.698	115.511	47.881	17. 554	52.330	16.308	38.448
MAMMA1000646	72.487	111.121	22.868	9, 213	27.074	81.604	46.859	34.04
MAMMA1000652	152, 920	94.568	319.943	76.610	67.817	87.605	41.747	77.720
		41.097	278.504	38. 131	36.289	67.327	34, 224	32.59
MAMMA 1 000657	116, 830						14. 102	35. 21
MAMMA1000664	48.908	37. 993	133.863	26.712	16.308	21.135		
MAMMA1000667	77. 285	24.312	99.732	25. 027	29.493	43.769	22. 193	24. 50
8330001 AMMAIL	42.561	28.100	-54, 970	17. 454	18.336	50. 398	38. 233	26.55
MAMMA 1000669	22.797	14. 382	57.803	14.670	6.337	12.841	7.392	12.08
			46.836	26. 498	25.826	33. 332	38.768	39.13
MAMMA 1000670	66.748	22.566			28.783	64.713	38.934	40, 59
MANNA 1000672	128, 331	25. 209	67.913	35. 262				
MAMMA1000681	66.397	40.677	32.249	14. 404	13.181	26.710	30.054	37.36
MAMMA1000684	85. 908	107.381	66.100	35. 992	32.881	41.006	36.719	77.83
MAMMA1000695	165. 293	107.442	551.458	130.714	88.510	70.985	43.857	55. 55
	82.316	25.689	52.797	22.639	22.884	48.899	39. 297	29.63
MANNA1000702					15. 562	39.909	33. 303	25. 37
MAMMA1000706	81.416	25.442	34.529	20. 432				
MAMMA1000707	128. 277	17. 100	51.835	15.001	33.473	48.628	46.555	24.07
MAMMA1000713	75.263	59.677	109.995	37.970	23.975	33.874	30.149	39.49
MAMMA1000714	228.366	288.017	246.261	56.045	25. 380	80.480	51.219	64.58
MAMMA1000718	98.208	92.149	245.750	79.940	49.064	50.180	40.223	49.03
					73.612	78.021	29.904	60.25
MAMMA 1000720	158.737	111.227	446.586	101.175				
MAMMA 1000723	64.930	49.053	148. 285	40. 276	28.806	19.434	18.845	24.78
MAMMA 1000731	31.516	11.357	68.834	12.436	11.755	7.989	7.536	7.36
MAMMA1000732	121.291	56.513	230.064	58.746	51.582	53.763	35.440	49. 33
MAMMA1000733	24. 525	14.171	58.717	16.852	7.153	14.100	8, 586	10, 63
			142.152	102.345	44.860	84.456	43.098	98.0
MAHMA 1000734	113.011	127, 466					82.810	69. 40
MAMMA 1000735	142.978	48.490	130.520	34. 595	40.252	73.418		
MAMMA1000738	110.304	61.504	28.831	38. 642	18.942	31.735	48. 926	35.1
MAMMA1000744	140.264	94.669	281.287	76. 261	79.000	63.977	43.557	40.3
HALMA1000746	26.385	50, 110	37.264	16.895	10.790	35.280	3. 177	111.0
		36.619	52.587	30.957	36.810	46.899	25.359	24.8
MAMMA1000748	73.879				28.537	43.075	32.581	61.0
MAMMA1000751	42.505	27.882	58.087	44. 924				44.3
MAMMA1000752	55.785	55.799	193.100		25.798	29.655	21.969	
MAMMA1000757	314.709	210.647	535.246		161.327		112.625	152.0
MAHMA1000760	218.937	178.377	534.346	131.736	100.173	95.443	58. 158	91.2
MAMMA1000761	147. 993	73.793	349.399		65.436	75.180	43.310	63.4
MANMA1000775	75.873	25.684	170.040			20.938	15. 825	18.9
						51.597	28.394	33.4
MAMMA1000776	101.206	81.986	253. 211			29, 701	17.866	26.4
MAMA 1000778	71.839	47.596						
MAMMA 1000781	67. 901	30, 437	97.580	126.658		29.056	17. 488	26.9
MAMMA 1000782	286.062	65, 796	174.951	84.753	88.062	151.891	90.446	86.3
MAMA1000784	135, 655	91. 366			65. 127	26.625	29.991	78. 5
	143.478						29.032	46. 2
WILLIAM TOUCH								
MAMMA1000788	62.822	41. 315						
MAMMA 1000798		86, 328					38. 532	
	132.633		1 A 4 A 7 1 5	99.538	80.592	81.887	42.150	57.8
MAMMA1000798 MAMMA1000802	132.633		372.241	1 33. 334				
MAMMA1000798 MAMMA1000802 MAMMA1000810	150.779	88.200					12.633	10.7
MAMMA 1000798 MAMMA 1000802 MAMMA 1000810 MAMMA 1000813	150.779 31.571	88.200 14.636	31.497	9. 531	9.356	14.627		
MAMMA1000798 MAMMA1000802 MAMMA1000810 MAMMA1000813 MAMMA1000814	150.779 31.571 197.602	88. 200 14. 636 134. 253	31. 497 279. 885	9. 531 107. 679	9.356	14.627 99.046	64.626	62.0
MANNA 1000798 MANNA 1000802 MANNA 1000810 MANNA 1000813 MANNA 1000814 MANNA 1000824	150. 779 31. 571 197. 602 65. 693	88. 200 14. 636 134. 253 21. 602	31.497 279.885 64.020	9. 531 107. 679 1 38. 421	9.356 82.142 35.405	14.627 99.046 29.268	64.626 31.671	62.0 38.
MAMMA1000798 MAMMA1000802 MAMMA1000810 MAMMA1000813 MAMMA1000814	150.779 31.571 197.602 65.693 146.098	88. 200 14. 636 134. 253 21. 602 70. 894	31. 497 279. 885 64, 020 157. 448	9.531 107.679 38.421 347.656	9.356 82.142 35.405 39.428	14.627 99.046 29.268 44.524	64. 626 31. 671 33. 051	62.0 38.
MANNA 1000798 MANNA 1000802 MANNA 1000810 MANNA 1000813 MANNA 1000814 MANNA 1000824	150. 779 31. 571 197. 602 65. 693	88. 200 14. 636 134. 253 21. 602 70. 894	31. 497 279. 885 64, 020 157. 448	9.531 107.679 38.421 347.656	9.356 82.142 35.405 39.428	14.627 99.046 29.258 44.524	64. 626 31. 671 33. 051	62.0 38.8 44.

Table 53

			F68 654 15	13 004 1	00 640 (	85. 964	57, 386	102.963
		138.792	503. 964 1	13.084	86.648 19.319	37, 537	13, 012	20, 655
AMMA1000841	44.843	37.288		28.351		78, 492	50. 804	35. 389
	74. 347	36,747	169.008	44. 926	48.610		5. 283	1.757
LAMMA1000843	8. 643	4.650	14. 084	4.758	2.185	8. 547	17. 728	15. 507
LAMMA1000845	40.044	33.955	33.012	21.488	15.747	23. 310		
	197.033	79. 321	307.054	96. 446	73.025	75. 853	98. 526 22. 320	72.039
MAMMA1000854	66.648	33. 221	63. 298	17. 429	20. 157	33. 288		21.685
MAMMA1000855	10.264	4. 185	17.702	3.794	3.995	2. 454	9, 158	3.568
MAHMA1000856	186, 269	40.945	84. 561	27. 973	38. 378	82.629	60.529	25.726
MAMMA1000859	64. 234	121.939	60.662	34.958	42.945	39. 557	20. 320	33.032
MAMMA1000862	40.107	21.345	23.693	16.808	28. 277	22.661	14. 238	14.693
MAHMA1000863	98.576	70.285	234. 996	67.796	55. 216	72.466	36.802	70.885
MAMMA1000865	1. 106	0.000	0.000	0.000	2. 321	0. 000 18. 758	0.000	0.000
MAMMA1000867	46.228	24.216	64. 376	21.736	17. 699		10.742	6.964
MAMMA1000875	124.814	80.537	231, 558	88.627	57.015	82. 859	46.826	53.611
MAMMA1000876	87.475	36.523	94. 191	19.763	21.466	42. 434	27. 201	24. 439
MAMMA1000877	201.968	107.716		164. 333	86.827	114. 380	80.171	97.872
MAMMA1000878	99.671	67.833	257.022	71. 323	29.066	47. 487	36.714	37.365
MAMMA1000880	76.396	60.884	153. 335	45. 836	17.649	44, 996	19. 238	15. 353
MAMMA1000881	63.646	33.072	177.731	43.034	30.410	31.086	12. 184	38.045
MANMA1000883	71.807	24, 931	43.109	16.630	18.675	40. 320	44. 419	55. 440
MANMA1000897	88. 466	0.000	7.404	0.000	0.000	0.000	0.000	0.721
8680001 PMILEM	380.818	62.977	134. 846	45.311	63. 221	164. 332	122.071	52.933
MAMMA1000905	97.555	63, 528	161.117	57.777	42.205	50. 312	28. 216	42.710
MAMMA 1 000906	57.788	33.146	125.096	29.019	13. 531	29. 380	16. 982	14.930
MAMMA1000908	30.597	19.222	40, 351	11.584	5. 445	10. 392	13.469	11.612
MAMMA1000911	9. 952	29. 425	3.998	9, 963	1.886	7.419	5. 350	126.406 18.859
MAMMA1000914	82.184	23. 137	69, 228	20.659	18.111	35. 329	22.616 47.680	26.801
MAMMA1000920	92.123	62.032	37. 206	16.675	15. 550	47. 235	35. 902	77.424
MAMMA1000921	107.169	69.026	207.821	102. 347	60. 403 40. 229	64. 787 51. 543	49.349	95.211
MAMMA1000931	211,796	140.234	424. 498	95.390	55. 532	61. 420	51.119	60.328
MAMMA1000940	145, 411	82. 982	268, 876	70.972	79.478	106.717	53.292	91.187
MAMMA1000941	182.800	134. 847	509.857	131.193	68. 234	90.801	63.506	75.814
MANMA1000942	195.078	123. 131	445. 428	117.435	89.006	81.092	51.063	85. 539
MAMMA1000943	196. 926	99.988		78.330	98.779	104. 172	79.021	96, 980
MAMMA1000952	161.019	97.081	355.265 14.918	11.103	5.840	41. 230	24.471	6.893
MAMMA1000956	43.741	16.217	225.645	64.794	42.610	47.323	34. 337	45.567
MAUMA1000957	95, 532	53.066	781, 968	204. 962	120.611	123. 900	84. 354	140.995
MAMMA 1000962	281.600	192.048	417. 591	111.282	64.746	81.685	51.694	78.953
MAMMA1000956 MAMMA1000958	151.087 217.975	107.043	313.251	58. 469	41.964	45.044	41. 392	63.998
MAMMA1000972	18. 150	48.148	119.482	22. 427	18.041	15. 672	12.870	33, 135
MAMMA1000972	36.667	18.879	24. 787	11.758	12. 527	19.441	17.828	22.312
MAMMA1000975	44. 972	19.058	38. 995	20.137	30.793	22.864	65.817	45.398
MAMMA1000976	122.625	67.075	216, 981	70.671	60, 470	91.475	60.614	81, 173
MANMA1000979	81.812	102.452	145, 415	68.435	53, 443	56. 902	38.749	89.759
MAMMA1000986	118. 211	39.368	239, 204	68.513	49, 208	56.431	42. 354	94. 152
MAMMA1000987	81.456	50.679	249.660	43.686	35, 580	49.753	23.004	41.997
MAMMA1000988	150. 907	68, 191	242.562	63.946	34.252	81.162	48.528	86.723
MANNA 1000994	101.984	21.000		21, 154	26.136	49. 152	44. 373	50. 523
	166,669	75. 193		91.202	105.673	107.213	56.957	84. 216
				47 070	24 216	35.674	26, 101	59.032
MANNA1000998			146.092	47.279	34.315	33.014		
MAMMA1001003	73.580	37.252			1.411	3, 633	0.800	
MANMA1001003 MANMA1001007	73. 580 3. 055	37.252 0.000	5. 547 65. 220	0.000 38,501				
MANMA1001003 MANMA1001007 MANMA1001008	73. 580 3. 055 40. 892	37.252	5. 547 65. 220	0.000 38,501	1,411	3, 633 18, 859 56, 674	0.800 47.979 44.237	31. 121 52. 328
MANNA 100 1003 MANNA 100 1007 MANNA 100 1008 MANNA 100 1013	73. 580 3. 055 40. 892 135. 486	37.252 0.000 31.048 126.855	5. 547 65. 220 372, 544	0.000 38.501 93.280	74.831	3, 633 38, 859 56, 674 20, 809	0.800 47.979 44.237 35.346	31. 121 52. 328 16. 624
MAMMA1001003 MAMMA1001007 MAMMA1001008 MAMMA1001013 MAMMA1001014	73.580 3.055 40.892 135.486 85.681	37. 252 0.000 31. 048 126. 855 25. 361	5. 547 65. 220 372. 544 77. 414	0.000 38.501 93.280 32.516	1.411 74.831 57.270 25.227 34.542	3, 633 38, 859 56, 674 20, 809 34, 975	0.800 47.979 44.237 35.346 35.352	31. 121 52. 328 16. 624 29. 728
MANNA 100 1003 MANNA 100 1007 MANNA 100 1008 MANNA 100 1013	73.580 3.055 40.892 135.486 85.681 93.867	37.252 0.000 31.048 126.855	5. 547 65. 220 372. 544 77. 414 180. 659	0.000 38.501 93.280 32.516 41.205	1.411 74.831 57.270 25.227 34.542 41.069	3, 633 38, 859 56, 674 20, 809 34, 975	0.800 47.979 44.237 35.346 35.352 41.726	31. 121 52. 328 16. 624 29. 726 36. 711
MARIATOD 1003 MARIATOD 1007 MARIATOD 1008 MARIATOD 1013 MARIATOD 1014 MARIATOD 1021 MARIATOD 1024	73.580 3.055 40.892 135.486 85.681 93.867 141.736	37. 252 0. 000 31. 048 126. 855 25. 361 49. 224	5.547 65.220 372.544 77.414 180.659 229.735	0.000 38.501 93.280 32.516 41.205 52.670	1.411 74.831 57.270 25.227 34.542 41.069	3, 633 38, 859 56, 674 20, 809 34, 975 54, 541 6, 698	0.800 47.979 44.237 35.346 35.352 41.726 4.305	31. 121 52. 328 16. 624 29. 726 36. 711 5. 091
MARIATOD 1003 MARIMATOD 1007 MARIMATOD 1008 MARIMATOD 1013 MARIMATOD 1012 MARIMATOD 1021 MARIMATOD 1024 MARIMATOD 1024 MARIMATOD 1025	73.580 3.055 40.892 135.486 85.681 93.867 141.736	37.252 0.000 31.048 126.855 25.361 49.224 49.918 8.964	5. 547 65. 220 372. 544 77. 414 180. 659 229. 735	0.000 38.501 93.280 32.516 41.205 52.670 5.843	1.411 74.831 57.270 25.227 34.542 41.069	3, 633 38, 859 56, 674 20, 809 34, 975 54, 541 6, 698	0.800 47.979 44.237 35.346 35.352 41.726	31. 121 52. 328 16. 624 29. 726 36. 711 5. 091
MARIATOD 1003 MARIATOD 1007 MARIATOD 1008 MARIATOD 1013 MARIATOD 1014 MARIATOD 1021 MARIATOD 1024 MARIATOD 1025 MARIATOD 1028	73. 580 3. 055 40. 892 135. 486 85. 681 93. 867 141. 736 13. 661 36. 353	37. 252 0. 000 31. 048 126. 855 25. 361 49. 224 49. 918 8. 964 24. 719	5. 547 65. 220 372. 544 77. 414 180. 659 229. 735 12. 310	0.000 38.501 93.280 32.516 41.205 52.670 5.843 10.363	1.411 74.831 57.270 25.227 34.542 41.069 13.733 34.518	3, 633 38, 859 56, 674 20, 809 34, 975 54, 541 6, 698	0.800 47.979 44.237 35.346 35.352 41.726 4.305	31. 121 52. 328 16. 624 29. 726 36. 711 5. 091 11. 316 23. 535
MARIA 100 1003 MARIA 100 1007 MARIA 100 1007 MARIA 100 1013 MARIA 100 1014 MARIA 100 1021 MARIA 100 1024 MARIA 100 1024 MARIA 100 1025 MARIA 100 1028 MARIA 100 1030	73.580 3.055 40.892 135.486 85.681 93.867 141.736	37. 252 0. 000 31. 048 126. 855 25. 361 49. 224 49. 918 8. 964 24. 719 27. 602	5. 547 65. 220 372. 544 77. 414 180. 659 229. 735 12. 310 14. 061 35. 295	0.000 38.501 93.280 32.516 41.205 52.670 5.843 10.363 20.296	1.411 74.831 57.270 25.227 34.542 41.069 13.733 34.518	3, 633 38, 859 56, 674 20, 809 34, 975 54, 541 6, 698 16, 233 14, 989 129, 655	0.800 47.979 44.237 35.346 35.352 41.726 4.305 15.746 25.031 96.375	31. 121 52. 328 16. 624 29. 726 36. 711 5. 091 11. 316 23. 538 134. 509
MARIATOD 1003 MARIATOD 1007 MARIATOD 1008 MARIATOD 1013 MARIATOD 1014 MARIATOD 1021 MARIATOD 1024 MARIATOD 1025 MARIATOD 1028	73.580 3.055 40.892 135.486 85.681 93.867 141.736 13.661 36.353 33.595	37. 252 0. 000 31. 048 126. 855 25. 361 49. 224 49. 918 8. 964 24. 719 27. 602 125. 555	5. 547 65. 220 372. 544 77. 414 180. 659 229. 735 12. 310 14. 061 35. 295 517. 898	0.000 38.501 93.280 32.516 41.205 52.670 5.843 10.363 20.296 181.208	1.411 74.831 57.270 25.227 34.542 41.069 13.733 34.518 15.861	3, 633 38, 859 56, 674 20, 809 34, 975 54, 541 6, 698 16, 233 14, 989 129, 655	0.800 47.979 44.237 35.346 35.352 41.726 4.305 15.746 25.031 96.375	40.973

Table 54

					** ***	AAA T	- 10- 100 T	c - A01	27 262 3
	MAMMA (001038	26.248	12.160	150.692	32.729	5. 309	10.436	15.903	27. 263
	MANMA 1001041	113.237	27.602	43.846	32.708	45. 924	47.820	46.929	16.614
5	MANNA 1001043	218.483	23.847	58. 163	22.306	10.449	41.046	45.779	31.087
Ş	MAMMA1001050	157, 361	80.096	220.216	71.548	69.197	49.684	13.493	49.872
	MAMMA 1001054	102.456	62.728	134.003	63. 324	43. 343	21.184	38.007	39.478
	MAMMA 1001059	136.357	48.942	59. 998	52.931	26.061	111.283	69.714	40.010
	MAMMA 1001066	387.798	103.377	293.890	140.850	119, 334	176, 295	158.563	60. 324
	MAMMA 1001067	82.327	39, 420	127.017	37.076	29.891	30.670	19.782	14. 257
		150.398	31,601	52.273	21.983	32, 143	57. 421	47.051	26. 375
10	MAMMA1001072			17.217	11. 406	43. 228	24, 053	24. 142	5. 176
	MAMMA1001073	101.957	23. 218	240. 332	94. 124	56.071	89.717	16. 387	14, 966
	MANNA1001074	104.201	41.827	23.705	29.782	21. 196	23. 184	14.757	17. 497
	MAMMA1001075	32.081	34.601		75. 869	35.841	49.660	67. 285	67. 244
	MAMMA1001078	102.185	111.402	317. 478			186.406	141.739	266. 507
	MAMMA1001080	367.248	210.764	130. 259	89.003 17.602	81.982 43.163	26.358	17. 452	14. 352
15	MAMMA1001082	50.264	39.773	20.039		18. 321	4, 593	0.000	0.000
,,	MAMMA 1001091	3.576	11.403	27.522	0.000			11, 524	
	MAMMA 1001092	50.554	25. 306	48.577	16. 425	T5, 153	18.849		4. 155
	MAMMA 1001094	353. 180	72.506	112.379	42.145	78.386	130.368	113.824	62. 964
	MAMMA 1001 105	138.777	111.2261	113. 121	82.426	80.960	45.158	16.891	45.652
	MAMMA1001110	15.141	8.661	7. 407	3.823	5. 537	6. 280	3. 216	4. 392
	MAMMA 1001126	299, 120	223.060	683.480	194.522	164.920	119.375	96.413	88.784
20	MAMMA 1001133	243.826	187.024	529. 603	144, 907	119.301	111.573	67.515	94. 605
	MAMMA1001139	291.212	867.784		473. 187	227.579	348.527	121.382	173.640
	MAMMA1001141	36.320	18.295	40.066	9.930	5. 202	26.277	16. 337	13.996
	MAMMA1001143	163.308	70. 387	153, 588	67.249	59.919	57.023	43.805	40. 903
	MAMMA1001145	110.718	43.148	141.067	30.89Q	31.851	11.000	10, 119	13. 322
	MAMMA1001150	80.076	29.005	50. 289	15. 249	7. 495	33.674	48.052	22.629
25	MAMMA1001154	203.206	129.777	429.878	121.700	90.014	77.333	45. 155	71.154
	MAMMA1001159	46.847	28.763	19.301	13.704	8.444	23.404	21.664	24. 248
	MANMA1001161	185, 601	233.229	485. 605	141.151	109.607	107.154	96. 161	79.043
	MAMMA1001162	196.299	51, 198	67. 587	29.962	40.684	78.949	43. 247	18.714
	MAMMA1001181	116.505	35.688	88. 127	33.728	40.701	41. 280	16.749	26. 312
	MAMMA 1001186	155.118	85.120	303.506	69.532	51.017	85. 295	42. 211	48. 082
30	MAMMA1001189	60.587	31.052	16.618	30.386	22.337	29.809	50.085	54. 044
• •	MAMMA1001191	120. 521	18.093	41.909	22. 249	21.661	39, 122	50, 157	24. 623
	MAMMA1001198	229.338	561.556	755. 924	695.028	205.811	536.623	412.766	746. 035
	MAMMA 1001202	322.950	274.854	664.569	248.672	218.550	168. 136	144. 829	179.567
	MAMMA1001203	170.551	101, 121	330.599	85. 243	72.915	53.390	44.564	52. 183
	MAMMA1001206	132.103	114.504	202. 256	65, 195	71.217	61.327	43.601	48. 988
25	MAMMA1001208	55.417	28. 101	30.608	21.282	25.686	27.394	20.016	15. 433
35	MAMMA1001215	199.721	123, 016	194.852	82,919	72.839	87.841	68.245	60.078
	MAMMA 1001220	223. 133	154.557	404. 346	110.968	91.387	74.073	58. 534	62.841
	MAMMA1001222	5. 585	4. 936	6.763	1.952	0.474	2.171	20.800	5. 022
	MAMMA1001223	94.809	29.294	42.345	15.601	20.861	20.316	32.446	15. 726
	MAMMA1001232	130, 199	45.692	227. 125	47.671	38.837	45, 692	59.906	32.862
	MAMMA 1001234	129. 344	27.935	227.692	95.815	64. 344	61.799	49. 210	34.673
40	MAMMA 1001237	29.560	11.083	23. 224	7. 241	4.489	20, 199	16.883	11.003
	MAMMA1001243	20.832	11.598	47.127	7. 253	32.689	20.073	7.954	6. 544
	MAMMA1001244	44. 925	10.751	11.473	9.770	11.102	14. 902	16.779	4. 470
	MAMMA1001249	43.758	23.671	15.616	19.023	10.556	26.846	10. 975	13. 758
	MAMMA1001256	169.303	81.917	266. 686	187.649	131.656	44, 850	55, 325	59. 786
	MAMMA1001259	70.213	24. 036	18. 445	18.447	25. 202	45. 289	34. 303	19. 546
45	MAMMA 1001260	154. 426	64. 153	81.115	52. 438	46.566	80.874	64.937	87.761
	MAMMA1001262	153. 326	53.618	54.054	40. 354	54.252	66.416	134, 449	25. 835
	MAMMA1001268	97.760	53.599	146.494	47.068	42.826	34.360	20.976	28. 286
	MAMMA1001271	305.116	66.364	106.518	32, 761	65.392	128. 314	130. 796	39. 913
	MAMMA1001274	73.329	94.857	235. 488	85, 814	64.385	71.860	51.097	62.114
	MAMMA1001280	65.399	17.595	13.218	9.853	3.831	37.015	12.303	6.374
50	MAMMA1001283	145. 535	67.060	129.301	56.055	38. 490	56. 397	52.661	34.076
	MAMMA1001284	253.434	60.199	204. 903	48.739	63.272	100.485	93.658	76.590
	MAMMA1001286	86.284	38.290	49.421	32. 175		57.666	59.470	32.210
	MAMMA1001289	169.737	90.053	62.200	32.142	102.670	66.398	64.913	47.082
	MAMMA1001292	103.898	20.400	28.796	15. 498	31.006	29.378	26, 545	31. 970
	MAMMA1001296	225.022	173,717	324. 251	133.662	60, 125	88. 173	70. 926	89. 316

464

Table 55

	DECEMBER 1885		- 00	****	30 705	10 040	75 575	27 020 1	27 020
	MAMMA1001298	80.875	60.189	230.669	38, 485	32.838	36.675	27.032	27. 836
	MAMMA1001305	153.258	67.563	147.529	35.286	31.766	65. 281	41.627	30.730
	MAMMA1001309	6.490	8. 306	6.534	3.627	4. 269	0.000	5.861	5. 705
5									
	MAMMA1001310	148. 253	53.093	165.786	46.753	41.171	63.488	82.639	54. 927
	MAMMA1001322	20.005	14. 809	29.403	19.332	11.227	14. 549	14. 163	15.700
	MAMMA1001324	82.605	28.652	85.996	52.506	31.339	47.688	30. 365	20.779
						15, 121			
	KANMA1001330	180.949	117.040	245. 119	52.680		97.891	81.121	27. 980
•	MAMMA1001333	101.707	75. 972	213.812	59.950	49.965	59.640	32.340	37.307
	MANUA 1001334	156,564	108, 340	81.315	64.901	34.949	73. 570	65. 555	73. 287
10						20, 426	44. 148	21. 930	33.068
	MAMMA1001337	105.507	35, 111	33. 563	17.119				
	MAMMA1001341	100.751	32. 100	79. 257	23.788	38.019	38.614	42. 286	29. 671
	MAMMA1001343	128.875	95. 425	301.822	74.316	77.337	85. 437	18, 963	98. 899
				40.648	21.963	23.320	30. 315	16.394	27.074
	MAMMA1001344	32.880	35. 930						
	MAMMA1001346	49.749	17. 537	51.635	21. 147	20.480	22. 107	26.805	24. 306
	MAMMA1001383	202.565	186. 453	597.532	117.676	100.238	103.083	68. 993	76. 274
15	MAMMA1001388	149, 105	66. 100	213.624	45. 488	52.686	66.868	85.346	57.974
	MAMMA1001396	197. 435	81.919	430, 433	80.848	94.812	95, 399	75. 293	90.889
•	MANMA1001397	116. 167	86.809	175. 125	67.323	58.676	56.833	61.558	52. 233
	MAMMA1001401	101.761	72.0901	194. 999	62, 960	48. 162	57.422	73. 403	78.023
					9.779	13.557	44. 301	11.008	20. 408
	MAMMA1001408	62.875	17.757	62.603					
	MAMMA1001411	271. 344	54. 507	67.489	20. 558	68.557	157.085	134.884	38.338
20	MAMMA1001414	74.836	21.511	88. 459	27.219	20.603	32.791	16.798	25.126
	MAMMA1001415	207.635	38. 228	51.690	26.716	68.700	89.184	99. 527	41.848
•						28.949		31. 339	23.195
	MAMMA1001418	103.090	35. 102	91. 976	39.234		27.016		
	MAMMA1001419	106.299	52. 357	210, 943	52. 570	45.256	41.351	37.624	25. 914
	MAMMA1001420	133.835	25. 587	149, 981	15.816	19.703	28.670	26.323	15.896
	MAMMA1001426		180.062	165, 308	87.320	89.096	170.869	109.848	84.772
		265. 539							
<i>25</i>	MAMMA1001428	310.313	180.134	229.960	136.337	147. 398	262.499	135. 345	83.047
	MAMMA1001432	266.375	107. 317	387.676	86.786	60. 159	83.974	37. 205	60.775
	MAMMA1001435	99.596	48.079	193, 151	53. 623	27.154	41.869	30. 388	39.835
					78.030		54. 359	43.154	50.728
	MAMMA1001442	103.071	100.872	193. 544		54.054			
	MAMMA1001446	180, 367	105, 551	197.748	98. 484	72.694	46.485	39.641	61.589
	MAMMA1001450	67.785	51.961	68.660	34, 362	32,591	32.211	28. 904	9. 424
	MAMMA1001452	180.732	124. 244	432.438	115.549	111.829	104. 153	92.517	96.081
30									
	MAMMA1001465	528. 588	255. 549	770.820	359. 206	364.762	388. 404	209. 219	264, 053
	MAMMA1001476	33.639	19.551	25. 289	5.909	17.988	24.584	26.252	17.981
	MAMMA1001478	117. 183	61.333	147.393	46.785	39.649	32.143	33.776	40.723
				31.646	28.808	44.671	62.901	69. 911	26.759
	MAMMA1001479	156, 131	59. 931						
	MAMMA1001487	67.613	53.042	92. 480	34. 978	30.928	40.427	27. 489	11.238
	MAMMA 100 1498	96. 522	111.213	222.159	50.813	14.811	23. 385	56.209	28.054
35	MAMMA1001501	216.969	55. 879	84. 459	38. 369	49.731	88.169	43.395	32.036
							54.854	34.762	36.860
	MAMMA1001502	124.674	57.815	131.281	46.452	43.478			
	NAMMA 100 1510	27.993	7.591	13.577	10.197	11.745	6.993	14.922	8.048
	MANMA1001522	56.601	24.819	109.236	27.569	21.472	26.994	29.481	17.416
	MANMA1001529	83.190	23.330	52. 489	20.883	31.879	41.170	29. 923	20.596
								25. 583	
	MAMMA 1001532	47.058	33. 575	98. 780	33.881	17.641	23.522		30.896
40	MAMMA1001533	97. 390	40.032	30.146	22.218	20.573	25.298	46.390	16. 233
	MAMMA1001534	0.341	0.000	0.000	0,000	0.608	5.274	0.000	0.000
	MANMA1001535	32.482	21.042	23. 902	24.788	14. 317	27.839	5. 277	10.537
	MAMMA 1001547	122.717	75.842	186. 325	45.519	45.073	43. 338	36.590	24.660
	MAIMA 1001551	103. 124	52.282	155, 615	43.540	38, 692	47.685	20.767	32.781
	MAMMA 1001569	47. 916	19.726	56. 549	24. 375	18.319	34.666	36, 128	11.381
15	MAMA1001575					29.095	50.896	55, 992	33, 156
45		137, 304	30.090	50.539	31. 981				
	MAMMA1001576	355, 571	57. 322	87.851	39, 259	62.142	115, 580	85. 589	39.636
	MAMMA1001584	59.860	30.398	60.438	23. 526	24.246	30.161	16.694	22.305
	MAMMA1001586	6. 157	32.887	0.000	2. 133	1.210	6,758	2. 949	4.371
	MAMMA1001590	150.616	76.439	214. 250	84.714	45.244	67.639	37. 913	52.869
	MAMMA1001599	40.717	29.889	37. 283	14.015	19.295	24.401	27.880	19.119
50	MAMMA 100 1600	109, 112	32.647	49. 324	13.148	24, 411	44.599	35. 258	20.344
50	MAMMA1001604					9.643	15. 339	24. 456	16. 253
		153, 185	34.765	63.275	52.861				
	MAMMA1001606	217.088	99.469	248.919	91.848	90.788	88.514	79. 192	78.377
	MAMMA 1001609	64,637	23.619	74. 281	18.302	10.063	9, 100	19,011	13.860
	MAMMA1001614	74. 839	29.828	9. 202	11.550	18.036	35. 992	21.716	14. 483
	MAMMA1001615	71.970	10.164	10.048	11.622	4.999	35.674	12.056	11.852

465

Table 56

				42 107 1	48 88 T	177 666 1	122 701	44.365
4AM4A1001619	361.714	65.104	138.945	35. 137	88.004	177.280	155. 721	
MAMMA 1001620	113, 233	68.799	320.014	88. 182	65.387	62.891	47.797	49.428
MAMMA1001623	32.719	15.493	22.246	8.396	13.561	16.233	7.490	7.940
MANA 1001626	75. 279	8.514	13.728	10.774	12.665	56.613	57.493	6.962
				8.734	4.064	8.190	14. 443	7.576
MAMMA1001627	28.468	7.652	39, 356					
MANMA1001630	36.419	36.649	115. 287	20.971	7.371	8.511	10. 371	16.570
MAMMA1001633	77.945	25. 597	143.786	22.273	51.279	40.689	37. 952	19. 350
MAMMA1001634	132.937	95, 570	297.140	83.974	56.835	62.263	58. 952	66.333
		47.359	225. 161	34.126	24.717	38.086	34. 792	34.698
MAMMA1001635	140.754				13.713	19.299	12.550	9.106
MAMMA1001649	30.569	12. 321	20. 513	11.727			62.712	58. 197
MAMMA 100 1654	150.282	91.691	90. 096	34. 969	64. 959	66.853		
MAMMA 100 1660	133, 470	97.805	42.199	61.020	54.089	65.813	66.019	54.874
MALMA 100 1663	394.964	202. 523	572.820	154. 372	162.177	148.843	118.542	79. 262
MANMA1001670	109.171	38.230	119.077	31. 362	18,030	43.797	53. 194	28.426
MAMA 1001671	145. 809	21.188	31.621	20.983	11.973	13.009	10.867	8.816
					8.375	23. 180	9. 271	18.786
MAMMA1001679	74. 490	17.313	20. 426	10.837				
MAMMA1001683	147.044	87.078	260.375	71.605	39.630	48.331	49.633	41.012
3831001AMMAM	12.824	14. 454	46. 223	12.860	21.575	12.528	5. 274	9.906
MAMMA 1001688	290.960	584.756 1	484. 182	407.752	105.060	319.616	241.392	1824.687
MANMA1001689	74.686	28.294	39.725	20.248	8.261	19.721	31.387	18.923
MAMMA1001692		64. 474	198.053	56. 976	35. 470	19.914	16.899	28. 825
	90.375				36.663	51.227	10.898	27.229
MAMMA 100 17 1 1	111.425	82.300	189. 195	30. 269			24. 223	13. 907
MAMMA1001715	67.545	40. 330	71. 553	28.616	19. 372	25.019		
MAMMA 100 1730	33.925	17.096	21.837	11.464	4. 477	36.743	11.375	8. 587
NAMMA1001735	79.384	42.172	38. 240	23.675	25, 390	20.932	27.963	11.313
MAMMA1001740	100.894	25.218	94, 454	17.836.	17.794	23.366	21.945	15. 107
MAMMA 1001743	199.112	118.364	141.535	72.049	46. 384	86.104	96.828	100.038
			0.000	2.086	2.551	2.098	5. 703	0.000
NAMINA 100 1744	23. 256	20. 454				125.697	46.388	55. 894
MAMMA 1001745	121.679	94. 047	301.292	106.455	100.677			
HAMMA 1001751	58.670	37. 967	90. 572	30.921	14.618	26.060	33.416	32.380
MAMMA1001752	284.221	89.024	175, 680	74.746	86.008	159.864	103.908	99.685
MAMMA 1001754	57.620	30, 193	53, 390	14.833	35. 182	39.454	17. 523	12.754
NAME A 100 1757	14.456	8.290	7.632	7.247	6.076	15, 580	5. 382	5.641
	283. 527	155. 103	596, 815	118.229	106.868	115.717	105. 154	147.707
MAMMA 100 1760					5. 043	11.697	22, 420	16. 539
MAMMA 1001764	33.825	15.661	33.885	14. 429				
HAMMA1001767	41.791	27.578	112.242	22.484	21.848	16.357	11.576	9. 367
MAMMA 1001768	50.861	34.645	129.707	25, 692	23.037	24.674	27.811	11.075
MAMMA1001769	206.737	82.818	645. 195	110.913	102.640	105.607	80.653	102.144
MAMMA1001771	123.973	30.551	49,772	16.877	55.099	52.348	41.113	48.806
MAMMA1001773	47.743	27.204	35.277	8, 450	18.002	17.141	23.713	30.755
	104. 585	49.519	92. 589	42.249	35.085	50.584	39. 215	26.862
MANMA 100 1778				82.231	85.003	87.248	61, 999	71.448
MAMMA 1001783	140.821	89.274	371.095				45.875	54.652
MAMMA1001785	119.072	65.819	256, 400	60.491	37.351	65.802		
MAMMA1001788	37.967	8. 305	25. 708	9.749	9.870	11.494	13.172	10.408
MAMMA1001790	202.092	181.258	279.482	57,700	22.737	29.284	28.819	46. 106
MAMMA1001800	24. 282	11.444	30.466	12.517	1.763	8.501	13.065	25.671
MAMMA 100 1804	150.744	16.771	51.213	14.975	33.630	67.533	64.799	20.701
MAMMA1001806	62.312	54.896	146.142	37, 371	11, 402	36.501	43.675	52.846
	17.002	11.569	32.023	10.166	5. 995	9.576	10.245	11.255
MAMMA 100 1812				19.704	15, 636	25.863	15. 187	22.130
MAMMA 1001815	50.743	27.272	61.778				7.974	11.601
MAMMA 1001817	10.653	7.578	15.446	7.044	7.758	3.611		
MAMMA1001818	48.733	19.657	87. 193	21,647	18,566	18.770	19. 255	18.678
MAMMA1001819	165.340	99.233	343. 318	111.523	112.261	57.848	73. 268	87.725
MAMMA1001820	48.662	22. 951	34, 879	16.243	11.743	9.468	15.897	11.396
HAMMA1001824	125.683	53.824	187.383	58.214	53. 591	47.999	45.347	37. 548
		30.370	42. 082		23. 518	23.996	20.045	8. 482
MAMMA 100 1832	56.633				43.465	44.952	56.814	24. 346
MAMMA 1001836	128. 477	58. 280	179.541	45. 913				
MAMMA1001837	118.428	66.031	172.658	60, 299	38. 153	37.090	17.947	50. 301
MAMMA 100 1848	42.562	27.622	82.759		20.435	22.941	15. 102	19, 124
MAMMA1001850	402.506	243.182	312.586	171, 182	143.034	232.615	91.466	106.637
MAMMA1001851	123. 305	30.035	69. 870	64.763	41.560	39.454	33. 329	45. 924
MAMMA 100 1852	198.774	161.311	321.896			112.820	91,724	
			234. 984			42.929	39.634	
MAMMA 100 1854	158.894	117.462						58. 279
MAMMA1001858	148.310	133.834	240.344	51.820	24.063	35.871	73. 151	1 30.219

Table 57

	F			*** ***	65 600 I	£0.00¢	67 000 1	CC 474 T	22 142
	MAXIMA 1001864	169.742	52.389	185, 785	37.880	50.896	67.999	55. 272	23. 142
	MAMMA1001868	82.643	56.419	59. 491	62.418	34. 438	47.003	29. 588	35. 585
	MAMMA1001874	9. 192	9.651	51, 178	7.405	11.275	9.054	7, 189	10. 453
5	HAMMA1001878	190.515	70.315	227.600	164.835	101.886	72.219	79.645	146.982
	MAMMA1001880	159.918	94, 489	292. 528	95. 467	48.528	98. 588	39. 271	81.114
	MAMMA1001885	117.729	44. 975	110.656	53.460	26.142	52, 223	41.423	29.156
	MAMMA1001890	127.969	47.712	247.654	60.558	29.367	36.838	39, 109	41.483
	MAMMA1001893	90. 120	22.271	50. 435	19.070	23.222	27.783	36,643	18.711
				188. 894	57. 356	38.856	45.633	22.050	26.367
10	MAMMA1001901	78.854	67.274			91.563	25.690	68.288	28.595
10	MAMMA1001907	159.767	70.062	305. 846	76.004		53.008	32, 123	40. 375
	MAMMA1001908	44.964	27.928	41.967	55. 852	40.219		0.000	
	MAMMA1001919	0.000	82.865	12, 109	0.000	2.270	0.000		5. 175
	MAMMA1001931	59.705	9.869	29. 213	49. 582	13.981	18. 165	29. 466	11.467
	MAMMA1001937	47.045	26.453	33, 302	16.535	17.844	31. 265	29, 899	19.650
	MAMMA1001951	114.033	76.574	311.618	70.531_	55.661	40. 552	39.990	40.224
15	MAMMA1001956	171.199	78.116	295.630	76. 171	65.654	47.426	<b>6</b> 7. <b>5</b> 68	57.411
	MAMMA1001957	114.304	40.789	155. 366	46.819	41.429	43. 671	26. 153	26. 982
•	MAMMA 1001960	99.822	63.449	192.955	55. 422	57.938	23. 395	42.027	44.844
	MAMMA 1001963	6. 938	3.651,	9.748	3.671	3. 137	0.000	0.000	5. 275
	MANNA1001969	237.109	164.919	517.768	178. 594	149.500	109. 284	97.612	137. 120
	MAJMA1001970	199. 358	123.085	297.080	101, 158	41.691	71.806	71.685	61.125
20	MAMMA1001978	1.206	0.000	0.000	0.000	1.081	1.561	0.000	0.000
40	MAMMA1001992	189. 502	91.630	283. 440	78.807	70.640	63. 218	71.282	32.898
	MAMMA1001994	85. 231	21.385	143. 259	40. 178	38. 484	54.686	24.893	33.837
	MAMMA1002008	66. 834	77.793	37. 647	14.813	20.016	33. 334	39. 365	10.388
	MAMMA1002009	144. 462	65,030	407.911	107. 350	55. 438	47.107	40, 434	57.138
	MAMMA1002011	32.832	13.901	27. 524	10. 188	19.701	17.344	22, 354	14.449
			67.057	159.576	65. 640	59. 239	37. 381	36. 122	50.747
25	MAMMA1002022	107.727	70.125	207. 390	72.614	55. 279	78. 953	108, 945	46.948
	MAMMA1002024 MAMMA1002032	176.885		362.313	98. 520	95. 826	104, 970	73.966	83.780
		270. 523	130, 983	303.660	81. 264	93.758	74, 391	34. 919	49.831
	MAMMA1002033	132.652	119.984			10.859	15, 705	11, 098	10.476
	MANNA1002041	19.611	15.313	18.901	14.070	30. 208	55. 486	24. 562	23.890
	MAMMA1002042	78.700	42.958		37.566	14.811	11.172	1.533	10.371
30	MAMMA1002045	7. 131	8.948	24.018	14. 459	45.781	34. 315	27.824	37.210
	MAMMA1002047	82.875	57.343	192. 240	55. 806		84, 218	104.806	75. 923
	MAMMA1002056	212. 189	152.323	474.785	146. 238	94.617	81.670	44. 227	65. 825
	MAMMA 1002058	149.112	126. 148	334.116	98.541	74.809	5. 917	16.902	5. 536
	MAMMA1002060	13. 278	7, 931	14.514	12.643	5. 782		63.052	39.667
	MAMMA1002065	128. 185	46. 405	127.810	82.855	59. 107	72.737		
0.5	MAMMA1002068	110.652	64. 982	163.753	51.583	45. 893	40.656	37.400	24. 128
35	MAUMA1002070	61.186	24.791	29. 988	16. 102	15. 306	31.362	22.002	21.338
	MAMMA1002078	170. 197	38.633	93.014	30.633	33.682	90.533	42.110	14.299
	MAMMA1002080	21.195	14.596	12.646	10.208	14.094	14.792	10.377	10.263
	MAMMA1002082	111.870	77.716	117.819	55.009	54. 940	28.457	25. 946	21. 254
	MANNA1002084	74. 297	40.086	152.790	30.118	30.052	28. 788	24. 428	24.140
	MAMMA1002087	17. 991	17.619	30.479	8. 932	13.026	13. 365	9. 996	6. 344
40	MAMMA1002091	78.604	26.611	41.258	17.086	26.812	39.757	46.803	27.660
	MAMMA1002093	17. 498	0.000	5. 942	5. 592	5.630	8. 103	11.278	4.689
	MAMMA1002095	78.790	13.430	22, 728	13.058	20.650	32. 157	32.621	8, 152
	MAMMA1002108	91.919	6.035	31.027	13.639	7. 939	32, 486	27. 923	11. 735
	MAMMA1002112	24. 376	27.337	10.667	11.574	. 5. 250	15, 678	14. 329	37.463
	MAMMA1002118	12.060	5.100	8.756	5.943	6. 502	7.856	7.396	3.149
45	MAMMA1002119	122. 271	36,908	59. 513	20.581	36.895	38. 172	39.046	32.476
• •	MAMMA1002125	159, 277	83.844	373.786	60.523	54.991	63.367	35, 366	35.797
	NAMMA 1002126	231. 380	139.298	431.047	153. 496	117.027	84.728	70.558	62.381
	MAMMA 1002128	102.647	35.864	48.863	19.098	20. 911	44. 235	39. 193	25.406
	MAMMA1002132	226. 752	118, 230	198.712	79.589	88.860	84.266	50.630	48.550
	MAMMA1002140	54. 642	53.227	115. 593		33. 524	31.026	24.905	32121
	MAMMA1002142	121.646	33.612	49. 214		27.295	103.698	68.348	39.850
50	MAMMA1002143	150.595	15.368	78.681	38.118		13.974	10.806	45. 937
	MAMMA1002145	237. 202	72.397	165.166		53. 986	87.872	73.605	22.437
	MAMMA1002147	73.366	34.088	45.076		33.648	53. 571	33, 082	8.766
	MAMMA1002153	133. 485	74.073				85.911	25. 126	19.099
	MAMMA1002155	320. 181	146.275				124. 338	93. 185	96.378

467

Table 58

								0.664
AMMA1002156	3.612	2.088	14.013	0.880	0.000	0.000	0.000	0. \$54
IANHA 1002158	70.916	40.655	88.575	24. 125	14.786	21.144	21.721	31.526
UNINA 1002 154	109. 211	29.584	54, 163	32.089	28.633	66.844	29.378	23. 138
AMMA 1002155		111.787		73.710	65.970	77.137	88. 540	53. 125
	166.029			1, 159	0.000	0.000	0.000	0.000
MMA1002170	0.000	0.000	0.000				87.695	71.650
IAMMA 1002174	139.902	178. 299		82.252	147. 225	141. 399		
MAMMA 1002175	49.635	20.661	21.290	16.108	13.918	22.449	12.875	18.564
AMA 1002180	117.470	55. 089	69.154	18.969	36.764	45. 946	59.721	45. 237
MAMMA1002198	123. 227	67. 539	235. 488	54.699	51.835	48.796	31.324	62.413
			420, 688	47. 331	61.775	61.499	42.296	74.029
MANMA 1002205	114.861	63. 437			32. 139	63. 320	64. 272	56. 392
MAMMA1002206	86.539	30, 685		17.788				
MANIMA 1002209	124.961	73. 557	143.211	32.601	43. 486	64. 448	43.661	36.987
MAMMA1002215	446.836	148, 590	401.477	50.983	162.248	310.059	210.563	225.764
MAMMA1002219	103.054	68. 338	110.047	29, 595	35.094	50.008	34. 183	47.670
MAMMA1002224	155. 329	135, 036	325.596	92.243	139. 113	54.888	50.692	104.338
				8.408	18. 280	19.024	14.880	18, 482
MAMMA1002229	54.055	19. 297	24. 594			62. 315	35. 205	65. 871
MAMMA 1002230	131.172	96.706	345. 936	76.632	50. 164			
MAMMA 1002233	40. 299	20. 503	27.780	14.645	13.380	24. 157	18.866	16. 294
MANMA 1002234	16. 951	13.815,	19.460	7. 251	4, 128	10.631	13.812	19. 438
MAMMA1002236	50.642	23. 553	50.683	14, 162	51.817	24.897	29. 324	44. 837
	88. 955	30. 943	38. 127	25. 451	21.889	37. 268	32. 369	10.849
MANMA 1002243			171.031	56. 513	74.300	48.863	11.431	66.114
MAMMA 1002250	101.569	23. 851					217. 429	157. 156
MAMMA 1002253	515. 165	161.871	322.750	80.630	175.660	370.878		
MAMMA 1002267	129. 167	239. 800	180.046	95. 357	56.654	98. 387	72.076	331.998
MAMMA 1002268	36.456	16.771	39.216	17.501	24. 043	16.873	20.704	13.929
MAMMA1002269	27.848	6.625	13.419	16.093	10. 154	9.666	6.915	4.635
MANMA 1002282	53.648	58.269	178. 298	38. 160	60.059	34.106	22.977	37.892
	62.491	17.873	48. 526	22.803	16.647	14.012	30.027	30.270
MANDIA 1002292				154.526	85. 449	104.060	60.152	54.729
MAMMA 1002293	236.280	162.513	481.000				25. 143	19.816
MAMMA 1002294	110.705	24.664	124. 002	36.492	33, 138	43.853		
MAMMA 1002297	65.424	40.774	88. 229	32. 940	16. 126	21.061	14.524	17.505
MANMA 1002298	104.368	30.772	64. 493	24.071	29.853	40.308	35.653	29.912
MANMA1002299	102.764	41.185	67. 139	29.656	30.944	33.813	19.722	23.248
MAMMA 1002308	69. 299	30.798	86. 503	30.668	29.756	27.771	17.935	16.223
MAMMA1002310	494. 257	272.509		186, 568	219.463	344.867	183.571	203. 149
		60. 941	315.707	69. 190	66.700	63.609	50.563	40.723
MAMMA1002311	151.653				19.878	36.852	19, 114	16.993
MAMMA 1002312	79.548	36. 483	113.839	34, 110				20. 920
MAMMA1002317	96.094	32.026	188.632	45. 170	45.365	46. 409	41.391	
MAMMA 1002319	141.320	69.599	218.472	74. 218	50.463	59. 927	44. 261	42.418
MAMMA 1002322	144. 393	65.401	253.730	67.857	46.931	25. 375	51.002	44. 820
MAMMA1002329	49.002	17.163	28.349	17.067	21. 239	27.218	20. 223	13.61
MAMMA 1002332	55.840	30.915	137.766	47.492	35. 312	32.956	23.130	16.41
MAUMA 1002333	75.478	17, 882	32.309	19.280	28.576	31, 145	41.629	17.63
MAMMA 1002335	171.866	50.373	149. 587	54.778	40.367	18.695	38.972	26.410
				63.915	53.097	48.035	33. 591	31.79
MAMMA1002339	91.741	62.618	152.049				45, 235	27.50
MAMMA 1002347	98.915	55.800	120. 784	40.650	55. 929	33.327		
MAMMA 1002351	70.045	22.016	35.600	18, 333	20.122	33.583	21.722	19.63
MAMMA 1002352	52.143	17.786	22.690	23.069	12.412	24. 411	13.818	11.94
MAMMA 1002353	128.336	52.785	144, 030	46.481	46. 561	36.806	12. 132	34, 57
MAMMA 1002355	46. 995	34, 505	123.684	29.737	22. 025	29.352	6.766	22.66
			86. 932	22, 189	25. 451	22. 826	13. 215	18.95
		1 21 772						
MAMMA 1002356	40, 901	21.732			142 084	59 794	89 656	1 51.18
MAMMA 1002356 MAMMA 1002359	40. 901 276. 825	92.529	330.418	168.428	142.084	59.794	89.656	
MAMMA 1002356 MAMMA 1002359 MAMMA 1002360	40.901 276.825 42.725	92. 529 25. 740	330. 418 47. 382	16.661	18. 409	9. 982	9.481	12.12
MAMMA 1002356 MAMMA 1002359 MAMMA 1002360 MAMMA 1002361	40. 901 276. 825 42. 725 152. 118	92. 529 25. 740 88. 131	330. 418 47. 382 201. 317	168.428 16.661 50.907	18.409	9. 982 51. 778	9. 481 26. 886	12.12 27.24
MAMMA 1002356 MAMMA 1002359 MAMMA 1002360	40.901 276.825 42.725 152.118 39.281	92. 529 25. 740	330. 418 47. 382 201. 317 119. 094	168. 428 16. 661 50. 907 21. 154	18. 409 41. 767 14. 517	9, 982 51,778 23, 579	9. 481 26. 886 14. 318	12.12 27.24 19.5
MAMMA 1002356 MAMMA 1002359 MAMMA 1002360 MAMMA 1002361	40. 901 276. 825 42. 725 152. 118	92. 529 25. 740 88. 131	330. 418 47. 382 201. 317 119. 094	168.428 16.661 50.907	18. 409 41. 767 14. 517	9. 982 51. 778 23. 579 65. 479	9, 481 26, 886 14, 318 60, 201	12.12 27.24 19.59 210.78
MAAMA 1002356 MAAMA 1002359 MAAMA 1002360 MAAMA 1002361 MAAMA 1002362 MAAMA 1002367	40.901 276.825 42.725 152.118 39.281 142.262	92. 529 25. 740 88. 131 22. 692 75. 867	330. 418 47. 382 201. 317 119. 094 50. 909	168. 428 16. 661 50. 907 21. 154 48. 285	18. 409 41. 767 14. 517 31. 065	9, 982 51,778 23, 579	9. 481 26. 886 14. 318	12. 17 27. 24 19. 59 210. 78 49. 29
MAMMA 1002356  MAMMA 1002359  MAMMA 1002360  MAMMA 1002361  MAMMA 1002362  MAMMA 1002367  MAMMA 1002371	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755	92. 529 25. 740 88. 131 22. 692 75. 867 66. 644	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090	168. 428 16. 661 50. 907 21. 154 48. 285 138. 658	18. 409 41. 767 14. 517 31. 065 42. 317	9. 982 51. 778 23. 579 65. 479 49. 599	9. 481 26. 886 14. 318 60. 201 32. 494	12.17 27.24 19.5 210.76 49.2
MAMMA 1002356 MAMMA 1002359 MAMMA 1002360 MAMMA 1002361 MAMMA 1002367 MAMMA 1002367 MAMMA 1002371 MAMMA 1002371	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755 90. 587	92.529 25.740 88.131 22.692 75.867 66.644 47.691	330.418 47.382 201.317 119.094 50.909 278.090 161.106	168. 428 16. 661 50. 907 21. 154 48. 285 138. 658 38. 559	18. 409 41. 767 14. 517 31. 065 42. 317 31. 139	9. 982 51.778 23. 579 65. 479 49. 599 36. 350	9. 481 26. 886 14. 318 60. 201 32. 494 34. 696	12. 17 27. 24 19. 5 210. 7 49. 2 25. 2
MAJMA 1002355 MAJMA 1002359 MAJMA 1002360 MAJMA 1002361 MAJMA 1002367 MAJMA 1002367 MAJMA 1002371 MAJMA 1002384	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755 90. 587 90. 935	92.529 25.740 88.131 22.692 75.867 66.644 47.691 85.538	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090 161. 106 249. 278	168. 428 16. 661 50. 907 21. 154 48. 285 138. 658 38. 559 71. 113	18. 409 41. 767 14. 517 31. 065 42. 317 31. 139 46. 508	9. 982 51. 778 23. 579 65. 479 49. 599 36. 350 40. 126	9, 481 26, 886 14, 318 60, 201 32, 494 34, 696 29, 975	12. 12 27. 24 19. 55 210. 78 49. 25 25. 27
MAJMIA 1 002356 MAJMIA 1 002359 MAJMIA 1 002367 MAJMIA 1 002361 MAJMIA 1 002367 MAJMIA 1 002367 MAJMIA 1 002380 MAJMIA 1 002384 MAJMIA 1 002384	40.901 276.825 42.725 152.118 39.281 142.262 119.755 90.587 90.935	92.529 25.740 88.131 22.692 75.867 66.644 47.691 85.538 7.306	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090 161. 106 249. 278 6. 051	168, 428 16, 661 50, 907 21, 154 48, 285 138, 658 38, 559 71, 113 7, 420	18. 409 41. 767 14. 517 31. 065 42. 317 31. 139 46. 508 . 3. 720	9. 982 51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699	9, 481 26, 886 14, 318 60, 201 32, 494 34, 696 29, 975 8, 116	12. 12 27. 24 19. 59 210. 78 49. 25 25. 22 44. 41 7. 60
MAJMA 1002355 MAJMA 1002359 MAJMA 1002360 MAJMA 1002361 MAJMA 1002367 MAJMA 1002367 MAJMA 1002371 MAJMA 1002384	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755 90. 587 90. 935	92. 529 25. 740 88. 131 22. 692 75. 867 66. 644 47. 691 85. 538 7. 306	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090 161. 106 249. 278 6. 051 66. 535	168. 428 16. 661 50. 907 21. 154 48. 285 138. 658 38. 559 71. 113 7. 420 12. 989	18.409 41.767 14.517 31.065 42.317 31.139 46.508 . 3.720 40.464	9. 982 51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699 53. 956	9, 481 26, 886 14, 318 60, 201 32, 494 34, 696 29, 975 8, 116 37, 080	12.12 27.24 19.59 210.78 49.25 25.22 44.41 7.60 19.51
MAJMIA 1 002356  MAJMIA 1 002359  MAJMIA 1 002361  MAJMIA 1 002361  MAJMIA 1 002362  MAJMIA 1 002367  MAJMIA 1 002384  MAJMIA 1 002384  MAJMIA 1 002385  MAJMIA 1 002385	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755 90. 587 90. 935 13. 712	92. 529 25. 740 88. 131 22. 692 75. 867 66. 644 47. 691 85. 538 7. 306 26. 458	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090 161. 106 249. 278 6. 051 66. 535	168. 428 16. 661 50. 907 21. 154 48. 285 138. 658 38. 559 71. 113 7. 420 12. 989	18.409 41.767 14.517 31.065 42.317 31.139 46.508 . 3.720 40.464	9. 982 51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699 53. 956	9, 481 26, 886 14, 318 60, 201 32, 494 34, 696 29, 975 8, 116 37, 080 20, 868	12.12 27.24 19.59 210.78 49.25 25.22 44.41 7.60 19.5
MAJMIA 1 00 2 3 5 6  MAJMIA 1 00 2 3 5 9  MAJMIA 1 00 2 3 5 9  MAJMIA 1 00 2 3 6 7  MAJMIA 1 00 2 3 6 7  MAJMIA 1 00 2 3 6 7  MAJMIA 1 00 2 3 8 4  MAJMIA 1 00 2 3 8 4  MAJMIA 1 00 2 3 8 9  MAJMIA 1 00 2 3 8 9  MAJMIA 1 00 2 3 9 9  MAJMIA 1 00 2 3 9 9  MAJMIA 1 00 2 3 9 9	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755 90. 587 90. 935 13. 712 119. 086 90. 573	92. 529 25. 740 88. 131 22. 692 75. 867 66. 644 47. 691 85. 538 7. 306 26. 458 32. 273	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090 161. 106 249. 278 6. 051 66. 535 97. 224	168, 428 16, 661 50, 907 21, 154 48, 285 138, 658 38, 559 71, 113 7, 420 12, 989 19, 547	18. 409 41. 767 14. 517 31. 065 42. 317 31. 139 46. 508 3. 720 40. 464 21. 438	9. 982 51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699 53. 956 26. 503	9, 481 26, 886 14, 318 60, 201 32, 494 34, 696 29, 975 8, 116 37, 080 20, 868	12.12 27.24 19.59 210.78 49.25 25.22 44.41 7.60 19.51
MAJMIA 1 002356  MAJMIA 1 002359  MAJMIA 1 002361  MAJMIA 1 002361  MAJMIA 1 002362  MAJMIA 1 002367  MAJMIA 1 002384  MAJMIA 1 002384  MAJMIA 1 002385  MAJMIA 1 002385	40. 901 276. 825 42. 725 152. 118 39. 281 142. 262 119. 755 90. 587 90. 935 13. 712 119. 086 90. 573 167. 171	92. 529 25. 740 88. 131 22. 692 75. 867 66. 644 47. 691 85. 538 7. 306 26. 458 32. 273	330. 418 47. 382 201. 317 119. 094 50. 909 278. 090 161. 106 249. 278 6. 051 66. 535 97. 224 370. 476	168, 428 16, 661 50, 907 21, 154 48, 285 138, 658 38, 559 71, 113 7, 420 12, 989 19, 547	18. 409 41. 767 14. 517 31. 065 42. 317 31. 139 46. 508 3. 720 40. 464 21. 438 82. 112	9. 982 51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699 53. 956 26. 503 77. 745	9, 481 26, 886 14, 318 60, 201 32, 494 34, 696 29, 975 8, 116 37, 080 20, 868 28, 921	7. 60 19. 51 14. 25 53. 90

Table 59

						A 17 T		T	1101
	MANIMA 1002400	10.797	7.113	11.587	4.041	5.847	4.732	4. 516	4. 194
	MAMMA 1002409	93.810	75.886	50. 232	41.725	30.159	43.673	520.771	70. 327
				76. 973	23. 185	26.301	31,997	16.726	11. 902
5	MAMMA1002411	81.111	34.713						
5	MAMMA 1002413	199.066	68.034	377. 354	55. 454	56.059	50.318	26.763	38. 961
	HAMMA1002417	30.976	26, 195	58. 136	15. 593	17.649	14, 266	7.765	11. 383
			47.715	208.629	48. 123	38. 391	40.117	26.156	31. 585
	MAMMA1002427	87. 721							
	MANIMA 1002428	108.360	83.671	293.146	88. 263	84.156	51.786	57.518	57. 126
	MANNA 1002433	90.843	23.726	38. 263	19.586	19.565	44.397	36.529	25. 042
	MAMMA1002434	117, 152	72.024	272. 113	68.694	66.706	54.616	45, 191	46.511
10					22, 955	36. 351	49.598	42.676	12.897
10	MAMMA 1002446	102.855	36.748	90, 796					
	MAMMA 1002 447	77. 962	49. 457	171.445	42.653	21.446	36.510	25. 929	27. 967
	MANNA 1002454	314, 500	201.950	539.572	188.845	118.797	99.696	72.794	103.951
	MANNA 1002 46 1	204, 681	47, 899	153.652	28, 137	56.943	63.968	55. 245	48, 401
						31,969	67.395	41.920	28, 713
	MAMMA 1002463	130.489	40.148	72.561	25, 745				
	MAMMA 1002 464	94. 697	34.520	44. 484	18.573	24. 045	50.857	37. 103	17.415
15	MAMMA 1002466	27.080	25.120	36.208	16.549	16.920	44.337	37.029	13.891
	MANIMA 1002 470	66. 277	10.542	19, 623	14.778	9, 384	20.022	21.241	15, 324
									26. 386
	MAMMA1002475	35. 982	25,009	77.707	23.670	24.685	10.963	12. 591	
	MAMMA1002480	85. 342	48.419	144. 499	40.755	50.788	48. 101	35. 187	30.058
	MAMMA1002485	256. 024	56.235	75. 461	32.978	72.095	120.038	77.311	49.943
	MAMMA 1002 494	66.749	23. 381	164.418	25. 376	48. 947	43.136	11.733	14.401
							26. 950	19.794	5. 551
20	MAMMA1002498	58.012	20.346	24. 265	12.932	13.125			
	MAMMA 1002524	73.628	20.842	11.923	21.047	20. 268	27.749	12.366	14.645
	MAMMA 1002530	82.789	19.903	43,603	13. 551	9. 151	28.535	27.989	12.505
	MAMMA1002538	101. 182	27.725	28.460	21.181	31, 900	45, 529	26. 380	25.658
	/						23, 145	30.820	51. 328
	MAMMA1002545	131.415	100.020	322. 993	72.173	54. 265			
	MAMMA 1002554	51.033	30. 923	62.549	16. 548	18.644	38.344	32.052	17.411
05	MAMMA1002556	201.613	52.773	211.073	70.139	99. 337	37.921	45. 357	46. 536
25	MAMMA1002561	199.748	128,004	586.968	135.854	118.280	54.740	81.217	\$1.656
						36.930	27. 532	51.392	13.777
	MAMMA1002565	57.918	43.508	20.564	13. 434				
	MAMMA 1002566	29. 155	16.405	7.906	3.460	1.967	13.518	5.709	5.318
	MAMMA1002571	73.034	22. 187	37. 154	25.594	6.079	28.030	19.946	20.955
	MAMMA1002573	218.479	62.669	183. 544	61.350	46.029	113.781	65.617	60.521
						22.615	43.283	55. 199	26.452
30	MAMMA1002576	109.621	18.498	33.802	10.617				
•	MANMA 1002584	244. 457	197, 626	384.879	79. 185	103. 251	112.917	113. 914	151.642
	MAMMA 1002585	133.865	28.963	56.983	17. 186	16.306	13.727	51.687	25. 753
	MAMMA 1002 586	67.168	39.043	34.776	15.656	19, 252	29. 596	35, 555	19, 945
				26.638	16.923	18. 956	18. 249	16.364	12.591
	MAMMA 1002589	98. 120	25. 567						
	MAMMA 1002590	268. 175	57.804	202.329	36.276	77.487	180. 923	123.883	42.552
	MAMMA1002593	131. 425	64. 951	130. 257	54. 131	23.515	55. 983	37, 410	35.272
35	MAMMA1002597	76.091	50. 352	131.097	33.606	42.551	25. 425	36.396	34.764
		69. 190		59.324	58.225	35.339	68.531	47.164	70.246
	MAMMA1002598		45. 133					64.732	
	MAHMA1002603	122. 932	40. 124	155. 801	51. 386	48.672	98.075		66. 103
	MAMMA1002612	330.999	152, 583	441.574	105. 603	112.764	175. 106	98. 853	99.475
	MANNA1002617	363. 139	211.631	557.754	145. 485	146.260	203.052	110.009	118. 254
	MAMMA1002618	90. 423	66.208	129.807	53.454	46.096	53.758	43.899	55.854
40			14. 223	23. 292	10.350	14.540	15. 236	12.465	13.642
70	MAMMA1002619	34.076				43.508			
	MAMMA 1002622	112.756	. 60 209	263.518	46.461		41.984	32.044	52.630
			60.308					4	
	MAMMA 1002623	89.689	68.083	149.811	64.401	102.216	102.611	54. 682	73. 325
	MAMMA 1002623	89.689	68.083	149.811	64. 401			54. 682 38. 497	73. 325
	MAMMA1002623 MAMMA1002625	89.689 83.660	68. 083 44. 949	149.811 94.038	64. 401 26. 154	102.216 32.540	102.611 34.576	38. 497	28. 162
	MANMA 1002623 MANMA 1002625 MANMA 1002627	89.689 83.660 9.090	68. 083 44. 949 2. 616	149.811 94.038 7.631	64. 401 26. 154 2. 675	102.216 32.540 0.000	102.611 34.576 3.940	38. 497 7. 852	28. 162 8. 826
	MAMMA1002623 MAMMA1002625	89.689 83.660 9.090 111.050	68.083 44.949 2.616 96.279	94.038 7.631 397.433	64. 401 26. 154 2. 675 77. 573	102.216 32.540 0.000 45.933	102.611 34.576 3.940 89.752	38. 497 7. 852 53. 737	28. 162 8. 826 108. 399
45	MANMA 1002623 MANMA 1002625 MANMA 1002627	89.689 83.660 9.090	68. 083 44. 949 2. 616	149.811 94.038 7.631	64. 401 26. 154 2. 675 77. 573 6. 679	102.216 32.540 0.000 45.933 3.741	102.611 34.576 3.940 89.752 10.219	38. 497 7. 852 53. 737 10. 741	28. 162 8. 826 108. 399 11. 301
45	MANMA 1002623 MANMA 1002625 MANMA 1002627 MANMA 1002629 MANMA 1002631	89. 689 83. 660 9. 090 111. 050 50. 470	68.083 44.949 2.616 96.279 10.960	149.811 94.038 7.631 397.433 11.524	64. 401 26. 154 2. 675 77. 573	102.216 32.540 0.000 45.933 3.741	102.611 34.576 3.940 89.752	38. 497 7. 852 53. 737 10. 741 8. 681	28. 162 8. 826 108. 399 11. 301 32. 169
45	MANNA 1002623 MANNA 1002625 MANNA 1002627 MANNA 1002629 MANNA 1002633 MANNA 1002633	89.689 83.660 9.090 111.050 50.470 32.234	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386	149.811 94.038 7.631 397.433 11.524 37.729	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053	102, 216 32, 540 0, 000 45, 933 3, 741 9, 358	102.611 34.576 3.940 89.752 10.219 12.456	38. 497 7. 852 53. 737 10. 741 8. 681	28. 162 8. 826 108. 399 11. 301 32. 169
45	MANMA 1002623 MANMA 1002625 MANMA 1002627 MANMA 1002627 MANMA 1002633 MANMA 1002633	89.689 83.660 9.090 111.050 50.470 32.234 59.898	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529	149.811 94.038 7.631 397.433 11.524 37.729 142.123	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014	102.216 32.540 0.000 45.933 3.741 9.358 15.348	102.611 34.576 3.940 89.752 10.219 12.456 18.150	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608
45	MAMMA1002623 MAMMA1002625 WAMMA1002627 MAMMA1002629 MAMMA1002633 MAMMA1002633 MAMMA1002636 MAMMA1002637	89.689 83.660 9.090 111.050 50.470 32.234 59.898 58.583	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541	149.811 94.038 7.631 397.433 11.524 37.729 142.123 11.323	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892	102, 216 32, 540 0, 000 45, 933 3, 741 9, 358 15, 348 14, 789	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.069	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104
45	MANMA 1002623 MANMA 1002625 MANMA 1002627 MANMA 1002627 MANMA 1002633 MANMA 1002633	89.689 83.660 9.090 111.050 50.470 32.234 59.898	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529	149.811 94.038 7.631 397.433 11.524 37.729 142.123	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014	102.216 32.540 0.000 45.933 3.741 9.358 15.348	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.069	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809
45	MAMMA1002623 MAMMA1002625 WAMMA1002627 WAMMA1002631 MAMMA1002631 WAMMA1002636 WAMMA1002636 WAMMA1002636 WAMMA1002636	89.689 83.660 9.090 111.050 50.470 32.234 59.898 58.583 55.442	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770	149.811 94.038 7.631 397.433 11.524 37.729 142.123 11.323 36.308	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176	102, 216 32, 540 0, 000 45, 933 3, 741 9, 358 15, 348 14, 789	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.069	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104
	MAMMA1002623 MAMMA1002625 MAMMA1002627 MAMMA1002631 MAMMA1002631 MAMMA1002636 MAMMA1002636 MAMMA1002636 MAMMA1002636 MAMMA1002646 MAMMA1002646	89.669 83.660 9.090 111.050 50.470 32.234 59.898 58.583 55.442 49.661	68. 083 44, 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800	149.811 94.038 7.631 397.433 11.524 37.729 142.123 11.323 36.308 69.217	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621	102, 216 32, 540 0, 000 45, 933 3, 741 9, 358 15, 348 14, 789 15, 750 64, 730	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.069 18.816 39.438	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014
45 50	MANNA 1002623 WANNA 1002625 WANNA 1002627 MANNA 1002629 WANNA 1002633 MANNA 1002636 WANNA 1002636 WANNA 1002648 WANNA 1002648 WANNA 1002648	89.689 83.660 9.090 111.050 50.470 32.234 59.898 58.583 55.442 49.661	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800 6. 907	149.811 94.038 7.631 397.433 11.524 37.729 142.123 36.308 69.217 9.595	64. 401 25. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621 4. 820	102.216 32.540 0.000 45.933 3.741 9.358 15.348 14.789 15.750 64.730 3.958	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.069 18.816 39.438 6.140	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742 8. 225	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014 6. 042
	MANNA 1002623 MANNA 1002625 MANNA 1002627 MANNA 1002629 MANNA 1002633 MANNA 1002636 MANNA 1002636 MANNA 1002646 MANNA 1002650 MANNA 1002650	89. 689 83. 660 9. 090 111. 050 50. 470 32. 214 59. 898 58. 583 55. 442 49. 661 15. 384 61. 935	68. 083 44, 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800 6. 907 69. 556	149.811 94.038 7.631 397.433 11.524 37.729 142.123 11.323 36.308 69.217 9.595 44.994	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621 4. 820 60. 882	102.216 32.540 0.000 45.933 3.741 9.358 15.348 14.789 15.750 64.730 3.958 59.089	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.816 39.438 6.140 42.135	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742 8. 225 62. 414	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014 6. 042 54. 651
	MANNA 1002623 WANNA 1002625 WANNA 1002627 MANNA 1002629 WANNA 1002633 MANNA 1002636 WANNA 1002636 WANNA 1002648 WANNA 1002648 WANNA 1002648	89.689 83.660 9.090 111.050 50.470 32.234 59.898 58.583 55.442 49.661	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800 6. 907	149.811 94.038 7.631 397.433 11.524 37.729 142.123 36.308 69.217 9.595	64. 401 25. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621 4. 820	102.216 32.540 0.000 45.933 3.741 9.358 15.348 14.789 15.750 64.730 3.958 59.089 8.462	102. 611 34. 576 3. 940 89. 752 10. 219 12. 456 18. 150 18. 816 39. 438 6. 140 42. 135 23. 347	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742 8. 225 62. 414 10. 991	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014 6. 042 54. 651 22. 157
	MANNA1002623 MANNA1002625 MANNA1002627 MANNA1002629 MANNA1002633 MANNA1002636 MANNA1002636 MANNA1002648 MANNA1002648 MANNA1002652 MANNA1002655	89. 689 83. 660 9. 090 111. 050 50. 470 32. 214 59. 898 58. 583 55. 442 49. 661 15. 384 61. 935 49. 617	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800 6. 907 69. 556 25. 105	149.811 94.038 7.631 397.433 11.524 37.729 142.123 11.323 36.308 69.217 9.595 44.994	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621 4. 820 60. 882	102.216 32.540 0.000 45.933 3.741 9.358 15.348 14.789 15.750 64.730 3.958 59.089	102.611 34.576 3.940 89.752 10.219 12.456 18.150 18.816 39.438 6.140 42.135	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742 8. 225 62. 414	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014 6. 042 54. 651
	MAMMA 1002623 MAMMA 1002625 MAMMA 1002627 MAMMA 1002629 MAMMA 1002633 MAMMA 1002636 MAMMA 1002636 MAMMA 1002646 MAMMA 1002648 MAMMA 1002648 MAMMA 1002655 MAMMA 1002655 MAMMA 1002655	89. 689 83. 660 9. 090 111. 050 50. 470 32. 234 59. 898 58. 583 55. 442 49. 661 15. 384 61. 935 49. 617	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800 6. 907 69. 556 25. 105 44. 430	149.811 94.038 7.631 397.433 11.524 37.729 142.123 36.308 69.217 9.595 44.994 13.568 94.935	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621 4. 820 60. 882 11. 569 34. 850	102.216 32.540 0.000 45.933 3.741 9.358 15.348 14.789 15.750 64.730 3.958 59.089 8.462 32.770	102. 611 34. 576 3. 940 89. 752 10. 219 12. 456 18. 150 18. 816 39. 438 6. 140 42. 135 23. 347 58. 417	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742 8. 225 62. 414 10. 991 41. 476	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014 6. 042 54. 651 22. 157 39. 910
	MANNA1002623 MANNA1002625 MANNA1002627 MANNA1002629 MANNA1002633 MANNA1002636 MANNA1002636 MANNA1002648 MANNA1002648 MANNA1002652 MANNA1002655	89. 689 83. 660 9. 090 111. 050 50. 470 32. 214 59. 898 58. 583 55. 442 49. 661 15. 384 61. 935 49. 617	68. 083 44. 949 2. 616 96. 279 10. 960 20. 386 50. 529 21. 541 29. 770 48. 800 6. 907 69. 556 25. 105	149.811 94.038 7.631 397.433 11.524 37.729 142.123 11.323 36.308 69.217 9.595 44.994 13.568	64. 401 26. 154 2. 675 77. 573 6. 679 16. 053 25. 014 5. 892 23. 176 43. 621 4. 820 60. 882 11. 569 34. 850	102.216 32.540 0.000 45.933 3.741 9.358 15.348 14.789 15.750 64.730 3.958 59.089 8.462	102. 611 34. 576 3. 940 89. 752 10. 219 12. 456 18. 150 18. 816 39. 438 6. 140 42. 135 23. 347	38. 497 7. 852 53. 737 10. 741 8. 681 38. 018 26. 406 26. 997 38. 742 8. 225 62. 414 10. 991	28. 162 8. 826 108. 399 11. 301 32. 169 22. 608 22. 104 38. 809 48. 014 6. 042 54. 651 22. 157

469

Table 60

								70
MAMMA 1002673	94. 294	135. 347	302.435	85, 978	115.544	122. 876	58.765	72.402
	169.486	32.550	60.424	32.013	39. 987	86.564	80.699	45.058
MAMA1002685	25. 020	18.401	21.785	11.312	11.628	3. 402	5. 660	25.002
MANNA 1002692	7.274	9.361	3.697	10.386	2.003	4. 100	3.302	9.849
HAMMA1002693	66,711	52, 339	15.641	32.934	10.671	20, 167	32. 429	30.795
MAMMA1002698	39. 272	32.200	43,657	33. 153	4. 354	11.796	12, 328	34. 409
MANMA1002699	18.348	10.645	5. 272	3. 333	2.314	3. 625	12.679	6.883
MANSKA1002701	66, 193	107.821	326. 150	82.189	33, 993	57.919	29.820	56, 144
		119.730	163.846	75, 850	65. 245	76.116	103.624	109, 697
	232.250		359.100	105.535	79. 020	76.543	25. 135	51. 975
	128. 862	101.834			18. 857	25. 978	44. 085	47,001
MAMMA1002712	55, 151	50.304	36.811	8.507	13. 366	39, 383	49.740	33. 088
MAMMA1002716	32. 821	37.741	37.674	23.554		57, 925		76.576
MAMMA1002721	128.620	78.060	360.516	86.920	49. 826		48. 421	
MANMA1002723	67.425	45.775	59.116	53. 954	27.853	31,646	28.039	37.993
MAMMA 1002727	4, 194	5. 317	4.081	4, 586	3.879	1.679	6.885	6.203
MAMMA1002728	45. 508	63. 239	134. 784	49.369	17.238	32.733	26.228	67. 828
MAMMA1002742	486.871	191.088	183. 567	79.031	108.740	257. 374	156.771	126.280
MAMMA1002743	17.914	25, 779	65.317	19.354	14.843	12.214	24. 184	22. 277
MAMMA1002744	70.172	65. 184)	190.550	59.599	40.023	33. 273	23.675	53. 991
MAMMA1002746	14. 967	8. 271	6. 293	9.116	3. 957	9, 800	1.039	7.011
MAMMA 1002748	53. 355	180. 986	171.425	25. 271	3.510	13.742	11.775	23,747
MARMA1002754	64. 093	69, 489	189. 499	44.022	29.371	15,039	15.857	30, 299
MAMMA1002758	25.835	7. 240	9.756	5. 507	5. 640	9, 500	11,968	9. 173
		58. 122	104. 988	33.940	18.698	86.679	92.471	84.012
MAMMA1002762	65. 824			59, 465	52.006	47, 508	45.629	48. 337
MAMMA1002764	104. 828	95.058	295. 803			30. 254	22.519	36, 212
MAMMA1002765	81. 926	54. 425	185. 685	56.838	25.634			15. 797
MAMMA1002769	20.078	9.062	33. 997	9.878	15. 366	12. 293	19. 431	929.910
MAMMA1002771	92.652	248.038	91.136	106. 297	36. 324	95. 235	52.022	
MAMMA1002775	51.236	37.084	125. 540	30.088	37. 975	21. 242	25, 695	24. 387
MAMMA1002780	23, 190	24. 572	73.778	29.564	12.337	13. 199	6.027	19. 175
MAMMA1002782	76.728	28.066	76.753	28.366	26.053	26.045	13.885	33.944
MAMMA1002795	17.412	3. 178	14.907	9.264	2.359	6.615	10.186	19. 921
MANMA1002796	28.596	28.390	48.340	13.930	16.360	14. 274	13.494	19.709
MAMMA1002805	25, 198	16.430	30.126	13.856	9. 933	47.769	23.312	13.432
MAMMA1002806	84, 431	28.564	34. 957	32.528	49. 335	29.125	31.705	30.489
MAMMA1002807	64.374	42.471	124.050	39.454	51.288	34.538	23.265	46. 125
MAMMA1002814	28.078	31,573	133.666	36.466	14.707	19.459	22.590	33. 539
MALMA1002817	8,719	10.443	6. 527	4.036	1.155	2.240	8.038	11.128
MAMMA 1002820	15, 173	5, 049	24.747	14.605_	7.416	9.432	16.038	5. 11
MAMMA1002830	91, 438	212.662	185.761	75.492	49. 491	111.835	311.632	133. 137
MANMA1002833	90. 875	71.138	237.238	50.346	44. 689	47.222	25.094	46.080
MAMMA1002835	28. 488	23.244	28. 102	14. 935	9. 504	12.597	16.302	12.70
MAMMA1002838	84.752	56.692	166. 200	49.694	30, 237	32.930	11.628	26.41
		53.519	151.675	23.902	32.033	41.236	27.950	47. 22
MAMMA1002842	98.706	31.051	107.479	18. 190	24. 282	30. 456	19, 401	13.72
MAMMA1002843	76.343	139, 150	228.560	66.881	72. 282	201.758	152.946	94. 16
MAMMA1002844	311.853		16.258	13.028	3.642	8.306	5. 338	22.84
MAMMA1002845	4. 464	5.631	235, 780	167.148	50.200	178. 228	129.737	278.80
MAMMA1002857	77.604	209.913		523.500	84.144	532.413	382.518	1000.09
MANAGE 10000	113.809	319.730	662.654	38.305	26.112	45, 735	86.883	51.98
MAMMA1002858				1 30, 305	1 40.114	1 43.133	1 00.003	1 31.30
MAIMA1002853	108. 297	33. 190			01 774	AC EET	28 420	50 46
MAMMA1002853 MAMMA1002858	108. 297 65. 375	102.643	253.035	92.062	91.774	46.567	38. 439	
MAMMA 1002853 MAMMA 1002858 MAMMA 1002869	108. 297 65. 375 85. 453	102.643	253.035 80.058	92.062 19.164	22. 933	26.217	42.600	30.85
MAMMA1002853 MAMMA1002858 MAMMA1002869 MAMMA1002871	108. 297 65. 375 85. 453 28. 097	102.643 22.923 6.998	253.035 80.058 5.660	92.062 19.164 1.623	22. 933 3. 087	26, 217 7, 477	42, 600 5, 467	30.85 3.40
MAMA1002853 MAMA1002858 MAMA1002869 MAMA1002871 MAMA1002875	108. 297 65. 375 85. 453 28. 097 20. 954	102.643 22.923 5.998 16.542	253.035 80.058 5.650 18.160	92.062 19.164 1.623 22.628	22. 933 3. 087 23. 110	26.217 7.477 21.099	42,600 5,467 24,952	30. 85 3. 40 32. 94
MAMMA1002853 MAMMA1002858 MAMMA1002869 MAMMA1002871	108. 297 65. 375 85. 453 28. 097	102.643 22.923 6.998 16.542 14.773	253.035 80.058 5.660 18.160 9.446	92.062 19.164 1.623 22.628 6.359	22. 933 3. 087 23. 110 8. 506	26, 217 7, 477 21, 099 13, 275	42.600 5.467 24.952 30.077	30. 85 3. 40 32. 94 23. 10
MAMA1002853 MAMA1002858 MAMA1002869 MAMA1002871 MAMA1002875	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288	102.643 22.923 5.998 16.542	253.035 80.058 5.650 18.160	92.062 19.164 1.623 22.628	22. 933 3. 087 23. 110 8. 506 12. 813	26. 217 7. 477 21. 099 13. 275 15. 447	42.600 5.467 24.952 30.077 20.107	30. 85 3. 40 32. 94 23. 10 22. 35
MAMMA1002853 MAMMA1002858 MAMMA1002869 MAMMA1002871 MAMMA1002875 MAMMA1002879	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288	102.643 22.923 6.998 16.542 14.773	253.035 80.058 5.660 18.160 9.446	92.062 19.164 1.623 22.628 6.359	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378	26, 217 7, 477 21, 099 13, 275 15, 447 18, 964	42, 600 5, 467 24, 952 30, 077 20, 107 34, 053	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41
MAMA1002863 MAMA1002868 MAMA1002869 MAMA1002871 MAMA1002875 MAMA1002875 MAMA1002880 MAMA1002880	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225	102.643 22.923 6.998 16.542 14.773 35.830 55.154	253.035 80.058 5.660 18.160 9.446 71.009 238.977	92.062 19.164 1.623 22.628 6.359 12.119	22. 933 3. 087 23. 110 8. 506 12. 813	26.217 7.477 21.099 13.275 15.447 18.964 34.101	42.600 5.467 24.952 30.077 20.107	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41
MAINA 1002863 MAINA 1002868 MAINA 1002869 MAINA 1002871 MAINA 1002875 MAINA 1002875 MAINA 1002880 MAINA 1002881 MAINA 1002885	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225 87. 039	102.643 22.923 6.998 16.542 14.773 35.830 55.154 28.425	253.035 80.058 5.660 18.160 9.446 71.009 238.977 35.323	92.062 19.164 1.623 22.628 6.359 12.119 25.333 14.016	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378 29. 952	26, 217 7, 477 21, 099 13, 275 15, 447 18, 964	42, 600 5, 467 24, 952 30, 077 20, 107 34, 053	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41 26. 27
MANNA 1002863 MANNA 1002868 MANNA 1002869 MANNA 1002875 MANNA 1002875 MANNA 1002879 MANNA 1002880 MANNA 1002880 MANNA 1002885 MANNA 1002885	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225 87. 039 398. 174	102.643 22.923 6.998 16.542 14.773 35.830 55.154 28.425 39.003	253.035 80.058 5.650 18.160 9.446 71.009 238.977 35.323 88.206	92.062 19.164 1.623 22.628 6.359 12.119 25.333 14.016 52.831	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378	26.217 7.477 21.099 13.275 15.447 18.964 34.101	42.600 5.467 24.952 30.077 20.107 34.053 61.975 39.216	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41 26. 27 20. 56
MANNA1002863 MANNA1002868 MANNA1002869 MANNA1002875 MANNA1002875 MANNA1002880 MANNA1002880 MANNA1002881 MANNA1002886 MANNA1002886 MANNA1002886 MANNA1002886	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225 87. 039 398. 174 45. 505	102, 643 22, 923 6, 998 16, 542 14, 773 35, 830 55, 154 28, 425 39, 003 7, 809	253.035 80.058 5.660 18.160 9.446 71.009 238.977 35.323 88.206 7.548	92.062 19.164 1.623 22.628 6.359 12.119 25.333 14.016 52.831 7.024	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378 29. 952 26. 325 9. 968	26.217 7.477 21.099 13.275 15.447 18.964 34.101 197.562 8.271	42.600 5.467 24.952 30.077 20.107 34.053 61.975 39.216 13.675	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41 26. 27 20. 56 5. 11
MAINA 1002863 MAINA 1002868 MAINA 1002869 MAINA 1002875 MAINA 1002879 MAINA 1002880 MAINA 1002881 MAINA 1002881 MAINA 1002885 MAINA 1002885 MAINA 1002887	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225 87. 039 398. 174 45. 505 65. 426	102.643 22.923 6.998 16.542 14.773 35.830 55.154 28.425 39.003 7.809 61.707	253.035 80.058 5.660 18.160 9.446 71.009 238.977 35.323 88.206 7.548 153.034	92.062 19.164 1.623 22.628 6.359 12.119 25.333 14.016 52.831 7.024 36.444	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378 29. 952 26. 325 9. 968 19. 739	26.217 7.477 21.099 13.275 15.447 18.964 34.101 197.562 8.271 40.974	42.600 5.467 24.952 30.077 20.107 34.053 61.975 39.216 13.675 38.649	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41 26. 27 20. 56 5. 11 41. 02
MAMA1002863 MAMA1002868 MAMA1002869 MAMA1002871 MAMA1002879 MAMA1002880 MAMA1002881 MAMA1002881 MAMA1002886 MAMA1002886 MAMA1002886 MAMA1002886 MAMA1002886 MAMA1002886 MAMA1002886	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225 87. 039 398. 174 45. 505 65. 426 58. 445	102.643 22.923 6.998 16.542 14.773 35.830 55.154 28.425 39.003 7.809 61.707 53.672	253.035 80.058 5.660 18.160 9.446 71.009 238.977 35.323 88.206 7.548 153.034 210.646	92.062 19.164 1.623 22.628 6.359 12.119 25.333 14.016 52.831 7.024 36.444 36.086	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378 29. 952 26. 325 9. 968 19. 739 31. 508	26, 217 7, 477 21, 099 13, 275 15, 447 18, 964 34, 101 197, 562 8, 271 40, 974 36, 186	42.600 5.467 24.952 30.077 20.107 34.053 61.975 39.216 13.675 38.649	30. 85 3. 40 32. 94 23. 10 22. 35 52. 41 26. 27 20. 56 5. 11 41. 02 35. 74
MAINA 1002863 MAINA 1002868 MAINA 1002869 MAINA 1002875 MAINA 1002879 MAINA 1002880 MAINA 1002881 MAINA 1002881 MAINA 1002885 MAINA 1002885 MAINA 1002887	108. 297 65. 375 85. 453 28. 097 20. 954 33. 352 46. 288 57. 225 87. 039 398. 174 45. 505 65. 426	102.643 22.923 6.998 16.542 14.773 35.830 55.154 28.425 39.003 7.809 61.707 53.672	253.035 80.058 5.650 18.160 9.446 71.009 238.977 35.323 88.206 7.548 153.034 210.646 25.600	92.062 19.164 1.623 22.628 6.359 12.119 25.333 14.016 52.831 7.024 36.444 36.086	22. 933 3. 087 23. 110 8. 506 12. 813 27. 378 29. 952 26. 325 9. 968 19. 739 31. 508	26.217 7.477 21.099 13.275 15.447 18.964 34.101 197.562 8.271 40.974	42.600 5.467 24.952 30.077 20.107 34.053 61.975 39.216 13.675 38.649 13.729 20.585	58. 463 30. 855 3. 40 32. 94 23. 10 22. 35 52. 41 26. 27 20. 56 5. 11 41. 02 35. 74 11. 29 21. 93

## Table 61

	MAMMA 1002898	88. 538	24. 524	42.725	9.653	16.551	32, 137	42. 359	30.615
				72.714	28. 234	32.209	91.200	60, 899	51.358
	MAMMA 1002905	191.445	19. 095						26. 917
5	MANNA 1002906	92.692	27.862	53.273	26.259	34, 130	57, 141	67. 635	
3	MAMMA1002908	77.656	66.964	209.054	54. 014	54.429	43.639	58. 625	50. 901
	MANMA 1002909	157.128	123.626	654.652	152.777	89.304	83.884	61.550	89.879
	MANIMA 1002918	55. 362	26. 201	35. 298	14. 931	10.960	19.166	27.775	29.119
	MAMMA 1002925	50. 571	70.116	54. 395	18. 071	27.814	43.511	11.984	57.467
	MAUMA 1002926	105.041	221.644	119.112	66. 217	73.866	245, 600	218. 974	550. 265
				147.112	32. 243	19, 181	31.875	24. 598	46.379
10	MAMMA1002930	68.089	38.713			38, 050	97.677	156.876	119.279
	MAMMA1002937	207.866	61.711	89.764	38. 377		15. 165	14. 230	14.534
	MAMMA1002938	34.139	13.727	21.350	7. 309	10.152			
	MAMMA 1002941	18.884	30.845	50.805	19.591	7.699	16.322	11. 528	24.529
	MAMMA1002947	63.095	31.441	46.623	20.590	18.624	28. 594	29. 987	39.586
	MAMMA 1002964	43.981	37.785	133. 836	22. 173	11.661	25.346	15. 389	28.296
	MAMMA 1002967	37.974	16.689	23. 126	13, 527	10.863	35. <u>085</u>	22,091	25.886
15 .	MAMMA1002970	178.268	124. 368	533. 590	120.984	97.317	92.795	65.069	109.854
	MAMMA1002971	99. 466	79.461	50.710	19, 662	15.091	40,745	37.592	51.546
	MANMA1002972	83. 922	33. 377	50.911	16, 436	12.354	42.113	50, 137	45.819
	MANNA 1002 973	117.540	70. 913,	318.513	45.601	38.568	34.070	22.903	68.699
			204. 398	227. 280	56. 459	375.745	119.386	122.750	226.538
	MAMMA 1002 979	80.771				0.000	0.000	0.000	5.076
	MAMMA1002982	19. 895	9. 493	14. 202	6. 265		22.630	15.594	36.952
20	MAMMA1002987	65.397	50.918	156.507	28.514	30, 958			
	MAMMA 1003003	104.891	69.630	125.933	48.800	36. 915	48.025	45.716	47. 346
	MAMMA 1003004	41. 353	106, 059	274. 522	111.746	92.691	59.597	33.719	77.654
	MAMMA 1003007	20. 423	21, 289	75.498	16.044	8, 909	15.878	6. 947	15.193
	MAMMA 1003011	45.615	37.641	29,754	23.843	21. 157	33. 395	48. 907	39.054
	MAMMA 1003013	65. 088	58. 284	49.438	27. 289	18.877	31.768	67.950	59.419
25	MAMMA 1003015	36.817	29. 585	89.251	19.826	4.679	16.602	6. 959	10.432
2.5	MAMMA 1003019	10.026	30, 107	5. 244	7.467	2.375	6,403	3. 225	6. 184
	MAMMA1003020	48,046	31,761	50.515	13.842	17. 142	19.341	28. 497	20.218
	MAMMA 1003026	28, 646	14.274	3.514	8.603	5.518	9.838	11.161	6.781
	MALMA1003021	248.219	140. 526	311.997	98, 494	105, 194	112.752	66. 452	132.570
		47.072	27. 208	130. 132	44. 811	42.096	33.806	17.555	36.757
	MANNA 1003033		49. 560	45.025	30. 912	25. 924	64.046	42.175	56.246
30	MAMMA 1003035	102.528		98.219	37.555	17.115	27.935	9. 656	25.906
	MAMMA1003039	37. 382	19.822			84. 250	59. 989	42.107	100.448
	MAMMA 1003040	76.014	95.416	243.138	114. 795		25. 258	13.745	23.444
	MAMMA 1003044	79.444	46.915	90.545	40.709	21. 121		175. 219	122.400
	MAMMA 1003047	376.340	121.483	150.100	91.015	100, 397	168. 621		
	MAMMA 1003049	26.899	9, 631	9, 169	2.907	5. 679	12.149	5.016	10.003
0.5	MANMA 1003055	38.639	24.977	76.695	21.811	15.758	11.937	6.277	20.034
35	MAMMA 1003056	31.238	13.811	32.121	15. 345	7, 891	17.689	3. 176	18.147
	MAMMA 1003057	68. 258	35, 596	34.053	23.862	19.335	28. 373	32.521	36.634
	MAMMA 1003066	43.837	46.015	117.875	31.178	11.361	17.068	9.179	35.831
	WAMMA 1003075	16.366	6.334	32. 529	10.374	3.215	6.507	2.433	11.804
	MAMMA 1003089	49.867	51.500	220.715	36.189	24.057	14.625	14, 530	41.852
	MAMMA 1003092	22. 129	73, 102	15,615	27.304	11.693	9. 575	15.986	84.953
40	NAMMA 1003095	8.240	37.313	24.078	8, 354	10, 123	9.662	24.609	12.392
. •	MANSIA 1003099	44.094	27.545	96.117	16.060	12.184	15.519	4.930	23.720
	MAMMA 1003102	44, 491	18.730	31, 447	14.500	22, 389	16. 929	20.089	20.899
	MANMA 1003104	35, 977	19.146	34.647	14, 588	10.720	11, 459	11.385	18.999
	MAMMA1003113	41.697	21.092	30.337	15.635	14.764	14.690	17.723	23.810
	MAMMA1003113	20.042	39. 595	102.916	21, 241	15.167	17.921	20.876	26.563
				102. 332	12.486	8.002	12.295	13,773	22.285
45	MAMMA 1003127	57.961	27. 221			86.667	135. 209	95. 293	83.256
	MAMMA1003131	267.516	37.924	129. 263	66.563	00.001	14, 984	7, 633	17.269
	MAMMA1003135	22.855	14. 308	5.624		2.690			5.312
	MAMMA1003140	6.575	9, 140	33.040	4. 487	0.895	1. 900	5.064	
	MAMMA 1003146	14. 105	18.018	18.562	11.213	11.461	16.500	8.591	9.815
	MAMMA 1003150	311.806	87.992	58. 938	77.271	104.739	165. 139	115.042	46.945
50	MAMMA1003154	93.002	39, 912	37, 471	22.819	19.655	31.742	26. 299	27.565
50	MAMMA1003155	41.709	26.308	36.508	14, 326	18.674	30.842	23, 489	18.046
	MAMMA1003157	34.876	32, 317	147.845	12. 108	24.093	12.999	8.766	19.930
	MAMMA1003163	37.900	25. 138	29.052		20.826	32.639	35.893	33.749
	MAMMA 1003164	26.961	14.747			5.852	14.778	13.694	20.137
	MAMMA1003166	12.213	5. 478			1.781	3.094	8.412	7.640
	[=v==v1003100	14.413	J			<u> </u>			

471

Table 62

	F-15-14-14-1	41 105	17 111	26.284	14.271	7.540	17, 180	16.220	11.568
	N89N31000010	31.105	17. 113			16,740	25, 216	14.845	17.364
	NB9N31000016	63.431	16, 195	24.879	17.001	36, 473	36, 956	51.575	34.673
5	NB9N31000043	87. 438	35. 161	58. 144	20.813			166.654	74. 328
3	NB9N31000045	83.399	109. 448	62. 101	95.653	93.734	94.218		13. 189
	NB9N31000054	41.821	12.636	37.831	15.025	15.265	18.963	10.894	
	NB9N31000076	22.822	22. 709	57. 320	14.223	12.517	9.029	11.713	24. 494
	NB9N31000086	31.281	74. 504	22.561	29. 164	11.744	29.951	13.909	30.012
	NT2RM1000001	11.595	9, 900	11.540	4. 467	4.016	8.823	6.775	5.184
	NT2RM1000018	333. 185	68. 022	171.103	77.680	48.418		122.906	79.595
10	NT2RM1000032	37.506	9.768	23.088	9. 453	13.222	16.128	22.911	12.495
	NT2RM1000035	185. 573	46.513	81.354	55.890	39.845	82.885	74.450	52.553
	NT2RM1000037	185.843	60.878	116.479	50.830	36.658	98.591	49.882	54.356
	NT2RM1000039	228.804	172.849	444.715	104.606	82.108	214.282	139.766	101.078
	NT2RM1000042	55. 479	102.774	112.292	145. 900	52.898	89.445	80.537	184.618
	NT2RM1000055	1.083	0.593	0.000	0.000	0.252	0.000	5. 227	0.000
15	NT2RM1000059	212.057	100. 267	173.989	78.130	50.792	143.445	83.189	102.504
		11.755	9, 438	11, 334	1. 925	2,705	2. 434	25.015	10.555
	NT2RM1000062		42. 956	56. 248	29.740	56.820	57.974	42.112	65.531
	NT2RM1000065	153.505		7. 914	2.716	6.609	8. 275	11.533	13.605
	NT2RM1000066	26.794	6.5394		97. 378	24.665	45.008	74, 491	266. 252
	NT2RM1000071	42.919	125.091	61.623	4.022	2. 135	8.919	13.254	4. 329
	NT2RM1000080	12.803	1.714	1.023		117.874	205. 973	155. 085	102.325
20	NT2RM1000086	393, 857	146. 368	283.360	100.835		0.000	5. 579	17.636
	NT2RM1000092	12.949	18.015	4. 187	6. 602	2.600	0.000	0.000	0.655
	NT2RM1000118	0.000	0. 276	0,000	0.180	0.000		19.700	10.812
	NT2RM1000119	18.719	5.828	9, 051	5. 794	3.873	6.048	1.697	3.614
	NT2RM1000121	2. 231	0.000	7.566	3.177	3.735	3. 309	141,553	
	NT2RM1000122	309.647	84. 904	138. 129	58. 379	75.966	213.166		57.569
25	NT2RM1000127	14. 133	3.707	2.380	2.322	3,743	4.212	8. 594	5.786
	NT2RM1000131	1.661	1. 269	0.348	0.000	0.768	0.000	2.271	2. 221
	NT2RM1000132	10.432	7.649	9. 599	3, 479	7.287	11.592	13.046	10.752
	NT2RM1000153	39.773	9. 302	10.314	3. 465	4. 419	11.775	17. 131	12.503
	NT2RM1000184	85.966	171.937	58. 982	34. 486	22.674	51.668	129.969	177.417
	NT2RM1000186	2.149	4.607	0.000	0.000	1.586	1. 226	3.974	7.121
	NT2RM1000187	29. 354	12, 303	16.019	17.222	15.020	17.176	15. 232	18.703
30	NT2RM1000199	16.274	0.000	17, 316	6.834	4.725	5. 212	8.917	6.720
	NT2RM1000213	17. 361	14.639	43.481	9.904	8. 998	12. 127	6.422	10.141
	NT2RM1000215	8.787	10, 858	90.070	4.505	89. 435	12.158	6.380	7.453
	N72RM1000218	0,000	10, 196	7, 239	2.227	1.452	4.273	8. 324	4.445
	NT2RM1000224	35.730	65, 418	0.000	47.537	20, 172	44.102	26.563	63.368
	NT2RM1000236	52.706	47.803	20.481	19, 138	42.513	21.813	58.118	100.492
35	NT2RM1000242	0.000	0,000	0.000	0.000	0.000	0.000	0.000	0.000
	NT2RM1000244	13.988	12.654	6.957	9. 937	5,047	8.026	8,918	3.968
	NT2RM1000252	283.006	144. 306	358. 324	169.383	149, 200	192.609	174.288	239.093
	NT2RM1000256	284. 496	113.021	203.771	67.954	94.270	152.181	132.435	150.452
	NT2RM1000257	8. 203	8. 081	9, 713	9.716	0.000	5,002	7.893	7.694
	NT2RM1000257	548. 461	312.072	494.663	164, 454	249, 491	313.672	232.568	270.549
40	NT2RM1000269	9, 472	7, 461	6.606		8,876	5.844	16.818	6.933
40	NT2RM1000285	8. 917	1. 259	3. 857	2,440	2.317	4. 289	4, 982	5.727
		83. 425		29. 246		22,650	25. 350	34.266	157.515
	NT2RM1000272		19. 960			5. 201	25.896	29.976	17.270
	NT2RM1000273	27. 031				11.534	34, 723	32,623	85.440
	NT2RM1000274	42.234				7.361	10, 397	4.200	10.119
	NT28M1000280	14. 289				9.099	9. 454	2.185	1.092
45	NT2RM1000295	8, 249				6.023	14.825	6.206	14.221
	NT2RM1000300	41.252		132 503			155. 874		
	NT2RM1000304	130.855				77. 271	189.937	108, 461	113.313
	NT2RM1000314	255, 347				114.936			25. 194
	NT2RM1000318	4.007				0.836	6.124	14, 151	14. 353
	NT2RW1000335						3, 558		
r0	NT2RM1000341	41.219							
50	WY2RM1000350								
	NT2RM1000354	6.027						2.303	
	NT2RM1000355	74. 362							
	NT 2RM 1000361	16.299							
	NT2RM1000365			0.00	0.000	1, 447	0.000	0.000	0.000

472

Table 63

55

				10 003	20 761 1	22 004 1	61 670 1	30 1/7 1	21 524
	NT2RM1000372	93.583	9.616	49.097	28.761	33. 904	61.678	39. 147	31.524
	NY2RM1000377	42, 186	17.871	22.783	12.668	13.142	15. 587	18, 377	23.602
	NT2RM1000388	8.811	19.351	1.155	5. 242	0.780	5.795	6. 201	11.464
5									
•	NT2RM1000394	0.899	1.862	0.813	1.925	0.438	0.000	0.000	0.000
	NT2RM 1000399	1.641	5.386	0.000	2.270	0.570	0.319	2.023	1.257
	NY2RM1000407	69.180	19.536	39.379	6.299	21.106	27. 229	14, 102	13.378
									0.000
	NT2RM1000421	0.890	0.000	0.000	0.000	0. 456	0.150	0.000	
	NT2RM1000422	102.028	152, 115	200.732	297. 482	65. 137	134, 344	50. 452	241.878
	NT2RM1000430	16.769	3. 286	12.402	4.398	4, 506	12, 149	11.238	7.508
10							81.561	72.026	118.786
70	NT2RM1000462	167.815	117.695	165.008	52.828	65.795			
	MT2RM1000499	16.037	22.127	75.152	12.507	7.415	7. 335	41. 299	22.217
	NT2RM1000512	126.510	24.122	12.786	25.082	11.161	46.878	21.802	31.090
						6,756	27.934	11.081	10.474
	NT2RM1000519	7.852	28.718	9. 178	14.716				
	NT2RM1000527	29.692	15.338	24.471	17.418	45. 221	59, 291	31.450	14.020
	NT2RM1000539	14,790	19.300	31, 135	14.824	2.560	6.669	3.751	10.774
15	NT2RM1000542	118.560	38. 555	21.020	20.675	29.849	30, 176	22. 378	32.507
.5									
	NT 2RM1000553	37.329	18.841	47.329	24.533	23.901	33.590	34. 084	33.966
	NT2RM1000555	77.352	46.168	43.953	21.772	15.838	16.936	12.057	35.840
	NT28M1000558	55.132	15, 424	20, 508	7.987	7.249	8.886	23. 984	21.919
						8.871	14. 324	12.341	13.462
	NT2RM1000563	39.161	14.058	17.872	12.234				
	NT2RM1000566	3.172	7.323	0.000	2.755	1.243	3. 584	2. 944	4.754
20	NT2RM1000570	65.428	72.508	44.124	24. 498	15. 164	26.341	21.720	56.340
20				9.841	14. 197	7. 525	7. 964	16.668	9.893
	NT 2RM 1000571	20, 300	15.881						
	NT2RM1000574	45. 305	32.953	5.746	5. 977	1, 945	5.060	1.526	3.809
	NT2RM1000580	10.540	9. 295	12.139	8.734	2.114	6. 532	5. 687	7.120
	NT2RM1000620	11.778	12.782	21.632	15.504	5, 894	4. 488	3. 359	17.303
	NT2RM1000623	3.914	2.515	0.416	3, 125	0.251	0.715	0. 355	2.159
	NT2RM1000630	17,633	6,091	6. 532	3.910	2.095	8. 257	7.963	5.411
25	NT2RM1000633	5.563	70.230	93.799	22.316	42.967	24.174	6.091	43.328
							0.000	1.258	3.039
	NT2RM1000634	3.427	3.869	2. 248	1.997	0.487			
	NT2RN1000642	87.902	31.353	26.846	11.421	21.495	75.074	66. 152	42.393
	NT2RM1000647	45.410	65.742	56.619	55. 351	49.439	30.233	26.128	50.923
	NT2RM1000648	25.285	9. 969	8.914	5.538	3.383	6.085	5.045	5. 201
								13. 836	
00	NT2RM1000650	22.370	15.864	19.881	11.036	29.031	8. 360		11.166
30	NT2RM1000661	23.325	6.294	12.692	7.551	6.360	11.076	18.036	9.158
	NT2RM1000666	13.966	1.244	3. 221	1.629	1.543	4. 997	1.079	2.418
						1.159	1.973	0.824	6.789
	NT2RM1000669	7.339	9.184	2.145	1.453				
	NT2RM1000672	58.162	25. 532	15.778	9.171	22.446	58. 987	16.791	14.945
	NT2RM1000681	21,724	106.663	3.979	14.842	2.185	20. 284	16.034	21.688
	NT2RM1000691	4.381	9. 202	2.832	3. 483	1.268	0.878	2. 181	3.652
35	NT2RM1000698	31.943	17.379	9.609	16.495	5. 185	8.614	8.628	12.092
	NT2RM1000699	10.439	2.722	5, 406	4.115	3.535	6. 367	10.784	8.214
	NT2RM1000702	32,110	7.097	17.438	3.946	5.019	19.783	16. 192	9.778
				20.468	14.964	19.912	19. 806	20. 940	16.286
	NT2RM1000703	32.168	17, 962						
	NT2RM1000704	25.926	35.690	22. 230	11.998	15.536	38.075	52.384	26.689
	NT2RM1000725	12.567	91,681	3.742	1 10.735	0.262	10.694	14.773	17.602
40	NT2RM1000726	7.525	9.354	5, 608	7.297	2.528	3.884	3.237	8.489
40	NT2RM1000731	144, 609	19, 850	46.338	14, 141	85.767	40.231	32, 791	30,972
	NT2RM1000741	14. 291	4.715	6. 122	2.576	3. 554	8. 230	5. 265	7.328
	NT2RM1000742	30.801	9, 241	6.240	6.116	3.655	11.131	7.680	11.315
	NT2RM1000744	69.419	21.887	27. 283	15.799	11.433	38.093	24. 162	24.347
								8.974	11.118
	NT2RM1000746	12.863	7.631	12.042	6. 326	6, 665	9. 321		
45	NT2RM1000747	24. 565	39.958	11.215	5. 537	1.866	7.009	10.940	21.461
45	NT2RM1000752	13, 148	7.585	3.359	5.748	4.905	1.290	6, 516	8, 586
						31.430	63.425	41.576	22.788
	NT2RM1000767	146.795	35. 621	33.719	11. 495				
	NT2RH1000770	24. 395	7.712	21.569	11.954	11.449	9.412	14.053	17.537
	NT2RM1000772	2.148	5.100	1. 271	2.181	0.000	1.505	6. 132	3.034
	NT28M1000779	284.561	185. 275	301.250		150.250	196.541	146. 279	96.926
50	NT2RM1000780	9.227	9, 621	4. 260	6.864	3.591	4, 298	8.898	2.912
50	NT2RM1000781	0.000	0.000	4. 468	0.666	2. 562	3.064	2.407	2. 127
	NT2RM1000789	79,877	28. 387	74.545		28.956	35.852	51.230	46. 548
				34. 906		6.856	4, 436	8. 934	3.531
	NT2RM1000800	4.947	10.706						
	NT2RM1000802	209. 372	41.025	60.767	12.693	69.721	155.310	133. 291	27.049
	NT2RM1000811	0.000	0.807	0.000	3.615	0.593	0.000	1.896	1.921

## . Table 64

	LITOOU LOOGGE	FF 47.	04 000	40 700					
	NT2RM1000826	55. 971	29.000	28.733	20.800	12.255	7.195	28. 144	23.708
	NT2RM1000829	19.377	19. 978	34, 233	28.539	40.659	14. 500	22.956	26,065
	NT2RN1000831	92.244							
_			176. 233	212.504	115, 234	47.485	121.255	114.428	264.692
5	NT2RM1000833	20.877	17. 302	8.876	4.821	8.474	6.471	16.424	13, 119
	NT2RM1000834	7.920	13.742	7.973	9.896	4.809	8. 919		
								6.281	8.562
	NT2RM1000841	31.899	32. 922	28.948	39.736	19.743	24.819	26.306	46, 020
	NT2RN1000848	10, 486	17.213	11.047	9.143	7.207	4, 310	8.632	18.858
	NT2RM1000850	4, 705	2. 700	0.000	1.784	0.000	1. 597	2. 104	7. 243
	NT2RM1000852	27,699	10.440	14.655	3.679	11.796	13. 435	15. 920	11.316
10	NT2RM1000853	0.000	4. 915	0.000					
					1.897	0.000	0.000	19. 505	3.017
	NT2RM1000855	295.899	111.992	196.426	53. 443	65. 232	138. 673	132.776	97.678
	NT2RM1000857	419, 515	279. 225	710.235	153. 528	198.222	264. 575	140. 191	196. 436
	NT2RM1000858								
		450. 537	223.032	628.109	128. 574	92.997	272, 161	183.324	165.845
	NT2RN1000867	36, 148	35. 491	71.518	26. 137	22.828	37.510	46.674	48. 259
	NT2RW1000874	94.766	25. 329	40,690	15.917	33.235	69.767	75.898	
15									34. 795
15	NT2RM1000882	32.751	18.077	43.528	12.957	13.381	12. 209	10.357	22.709
	NT2RM1000883	312.282	118.317	233, 345	90. 226	109, 110	311, 111	130.746	182.823
	NT2RM1000885	252.089							
			146. 253	191, 597	129.087	63.370	152.039	156.686	193.445
	NT2RM1000893	28.474	12. 532,	13.539	21.087	13.367	23. 959	22.465	14, 066
	NT2RM1000894	246.338	100.240	188.863	51. 822	48.537	189.474	182.264	80.716
	NT2RM1000898	8,028	11.716	12, 431	3. 461	8.055	10.349	3.262	8.889
20	NT2RM1000899	20.978	2.796	3.034	4.018	6.936	7.286	6. 525	8.715
	NT2RM1000905	90.972	37.943	146.214	36. 300	72.541	61.959	55. 239	45. 935
	NT2RM1000910	21.235	22.607	15, 176	6.355	3,770	20. 204	15. 343	18.656
	NT2RM1000914	199,944	90.792	169.446	46. 593	65.449	122, 556	87.145	72.117
	NT2RM1000919	36.141	16. 161	19, 116	13. 229	8.891	18.002		
								10.279	10.389
	NT2RM1000921	0.242	1.831	11.629	2.787	0.000	1.344	1.305	2. 292
25	NT2RM1000922	13.119	18.060	5. \$55	12.140	3.037	3.684	6. 526	16.464
	NT2RM1000924	29.895	12.894						
				4. 946	4. 788	7. 984	10.841	16. 108	5.749
	NT2RM1000927	48.046	34.032	49. 155	23.882	14.687	14.867	17.603	20. 582
	NT2RM1000951	13.349	11.379	12.531	13.272	6.919	7.215	10.192	8.882
	NT2RM1000956								
		5. 337	16. 522	6,739	2.246	6.192	6.379	6.215	8.675
	NT2RM1000960	24.574	14.841	49. 930	16.747	44.584	52. 121	23. 270	34.312
30	NT2RM1000961	20.594	16.610	28. 449	33.770	11.295	30, 987	65.017	30. 389
30									
	NT2RM1000962	1.479	8. 158	49. 309	6.863	4. 421	9. 226	13.337	10.245
	NT2RM1000973	69. 241	51.561	16.390	19.560	15. 357	27.890	33,675	45.410
	NT2RM1000978	0.000	0.000	0.000	0.000	0.000	0.000		
								0.000	0.368
	NT2RM1000982	7.275	2.308	2. 120	2.059	1.138	1.293	1.746	4.769
	NT2RM1000991	13.759	6.798	22, 345	7.467	6.192	11, 494	1.942	9.039
	NT2RM1000994	12.087	15.119	14.969	10.866	9. 132	2.303		
35								4.549	14.654
	NT2RM1001002	46.263	5.707	19.271	15. 499	18.065	33. 283	21.225	33.831
	NT2RM1001003	14, 107	33.647	23.710	23.835	3.391	10,638	8, 307	14.681
	NT2RM1001008	4. 937	4.696	0.740	4. 466	2.544			
							3. 192	3.215	10.971
	NT2RM1001011	67.834	16.031	21.431	8. 274	20. 203	45.979	40.030	18.121
	NT2RM1001013	25. 323	6.694	3. 303	6.673	8.650	15.882	23.168	23. 126
	NT2RM1001017	8.644	4. 934	1.214	2. 455	1.873		4.062	
40							2.894		7.068
	NT2RM1001018	224.654	234,771	124.092	68.774	75.070	85.777	124.713	184.612
	NT2RM1001026	23.853	12.510	10. 387	14, 301	5. 568	12.341	14.618	17.008
	NT2RM1001028	11.717	13. 271	17.437	18.862	5.641	12.231	8.930	
									11.443
	NT2RM1001043	21.614	13, 830	4, 261	8, 481	4.770	7.687	17.274	10.663
	NT2RM1001044	21.983	20. 272	44. 315	8, 181	4. 171	5.809	4.623	9.566
	NT2RM1001059	3. 169	2.991	1.316	0.000				
45						0.352	2.127	2.878	3.632
,•	NT2RM1001063	0.879	5, 544	0.768	1, 254	0. 973	4, 181	1,761	5. 391
	NT2RM1001066	3.011	3.061	0.000	3. 241	• 0.000	1.348	1. 228	
	NT2RM1001072	13.706							3.011
			7, 601	5. 972	2.306	0.165	3, 139	5.672	5.851
	NT2RM1001074	32.455	14, 324	28.723	10.090	6.573	10.841	7, 837	10. 538
	MT2RM1001076	7.339	4, 891	0.792	2.511	0.000	5. 644	6.602	2.026
	MT2RM1001082								
50		63.705	50. 432	105. 417	34.113	20.331	17. 230	16.378	21.799
50	NT2RM 1001085	13. 921	7.236	4, 420	3.206	4. 563	0.966	5, 984	4.704
	NT2RM1001092	16. 133	28, 559	80. 293	36. 442	13.840	23. 671	15, 948	30. 844
	NT2RM1001102	2.299	0,000	0,000	0.000	0.000	2.006	1.301	2.772
	NT28W1001103	4.293	14, 550	11.888	3.980	17.852	8.345	2, 505	12. 387
	NT2RM1001105	0.000	0.418	0.000	0.686		0.000		
						0.000		0.000	1. 156
**	NT2RM1001112	6. 983	5, 403	12. 985	7.889	7,226	5. 412	8, 469	12.089
5 <i>5</i>									

Table 65

				Tal	ble 65				
	NT2RM1001115	100.486	24.788	67.251	18.301	19. 421	53, 304	29.318	21.097
	NT2RM1001122	18.980	19.515	19, 938	11.109	10. 211	34. 308	33. 955	13.422
	NT2RM1001136	4.811	3.751	2.520	1, 126	0.765	2.194	2.817	5.117
5	NT2RM1001139	78.791	18.931	27.710	8. 382	21.060	31.349	14.028	14. 521
	NT2RM2000003	27.773	13. 438-	12.296	3. 254	10. 288	4. 103	14.697	22.880
	NT2RM2000006	64.154	36.637	117.073	30.277	27.783	25. 842	17.647	24. 349
	NT2RM2000010	57.806	33. 217	60.148	20.749	86. 788	23. 487	19.722	22. 551 36. 344
	NT2RM2000013	24.877	27. 244	40.874	15.590	40.045	30. 831	48. 932	27. 939
10	NT2RM2000030	68.595	26.308	27. 271	17.595	26.608 13.094	11.681	12. 137	11.426
70	NT2RM2000032	22.984	13.418	59. 847 28. 101	11.737	9,740	51.053	23.006	23. 405
	NT2RM2000039	35.892	5. 887	20. 886	10,060	5. 098	11, 101	20, 459	10.744
	NT2RM2000042	7.936 12.085	9, 200 11, 085	15, 415	5.779	5. 195	6, 720	11, 106	5.712
	NT2RM2000092	51. 998	31. 271	57.365	24.041	26.832	24.640	12.930	20.135
	NT2RM2000093 NT2RM2000101	34, 341	45. 587	64. 294	27.692	29.563	48. 487	33. 388	54.246
15	NT2RM2000104	73, 163	48.315	58, 786	33, 739	39.845	53.753	69. 151	73.279
	NT2RM2000124	35, 818	16. 923	31, 954	10.723	11.012	23.770	21.401	72.254
	NT2RM2000155	31, 139	23.019	27,033	12.467	9. 797	13.085	10.315	17.050
	NT2RM2000191	151.075	54.651	87.171	59. 579	62.006	74. 514	126.950	91.326
	NT2RM2000192	0.760	2.690	0.971	4.582	1, 137	2. 242	1. 413	0.000
	NT2RM2000239	92.578	36.060	71.933	31, 157	21. 570	60. 155	49. 672	39. 127
20	NT2RM2000240	104,218	69.966	77. 545	23.453	53. 412	78. 029	64. 223	83.906
	NT2RM2000241	70. 281	31, 167	42.733	18.007	14. 544	13.466	26. 176	42. 298 50. 848
	NT2RM2000250	72.366	22.586	52. 512	23.631	19.076	29. 100 44. 083	50.616 74.558	29.086
	NT2RM2000259	90, 122	33.799	39.931	17.198	9.865 77.794	188.072	216.739	59, 426
	NT2RM2000260	340.036	40.469	141.952 38.440	1, 951	3, 495	14, 217	14. 995	14.683
25	NT2RM2000265	24.506 131.692	88.080	127.535	51.611	38. 294	53. 574	55, 104	70.583
	NT2RM2000306	45, 342	24, 950	44. 593	13.884	40, 471	40, 133	22.666	33. 254
	NT2RM2000312	13.383	57.043	78.915	13, 258	60.055	90. 975	183.675	38.391
	NT2RM2000322	33.318	18.077	22.354	11.030	6.002	8. 829	16.962	15. 344
	NT2RM2000343	70.618	78.514	302.242	43.179	64. 338	35. 838	84. 150	77. 161
	NY2RM2000359	79, 203	25.437	34. 945	19.556	16. 348	47. 922	31.041	20.663
30	NY2RM2000362	138.367	75.052	100.195	73.363	49. 276	128. 683	126.847	106.528
	NT2RM2000363	41.249	17.128	40.363	12.316	18.047	6. 982	11.907	9. 239 93. 203
	NT2RM2000368	225.356	121.451	100.718	49.727	89.663	128. 354 80. 287	136, 054 140, 890	131.756
	NT2RM2000371	88.897	208. 325	97. 848	212.525 34.316	33.081 25.750	36.072	34. 151	51.955
	NT2RM2000374	54. 398	55, 656 35, 012	153.004	24. 245	23.611	19.094	24. 288	26.745
35	NY2RM2000387 NY2RM2000393	31.537 43.873	18.662	32.917	12.496	14, 167	17.560	23.452	33.102
33	NT2RM2000395	11.936	2.901	3. 145	1.722	4.564	6. 102	4.725	9. 257
	NT2RM2000402	26. 540	28.616	42.681	18.209	10.970	24.876	<sub>~</sub> 20.077	26.993
	NT2RM2000405	29.390	26. 302	56.236	18. 191	18. 624	17.673	19. 408	19, 435
	NT2RM2000407	213.973	77.583	145. 459	42.798	73. 678	124. 360	103. 989	122.635
	NT2RM2000410	46.375	23.782	29.096	10.711	13. 131	26.855	27. 992	20.820
40	NT2RM2000420	41.781	29.100	39.676	24.872	16.605	26.730 186.812	29. 136 131. 563	43.708 125.088
	NT2RM2000422	400.274	145. 824	265.042	51.828	73. 571	60. 353	42. 529	86.903
	NT2RM2000423	119,707	56.563 24.735	36.727	13.780	10.160	32, 134	23. 468	26.716
	NT2RM2000452	28.062	19.762	14, 685	5. 603	7. 485	22.242	10.716	6.249
	NT2RM2000490	57.984	29.556	42.743	15.403	19.316	36.503	21.105	31.221
45	NT2RM2000497	44.862	39.966	107.651	23.488	15. 277	19.316	13.374	15.412
45	NT2RM2000502	49.184	33.683	39.515	14.256	18.792	23. 598	23.921	27.178
	NT2RM2000504	53.653	30, 376	45.453	19.836	22.267	39, 106	28.508	19. 188
	NT2RM2000514	40.702	23.938	23.980		12.601	20. 319	19.147	27.441
	NT2RM2000522	6.782	0.000	4.730		1.616	2.008	4, 021	14. 506
	NT 2RM2000540	28.543	24. 938	24. 326		9.799	16.595	10, 471	17.045
50	NT2RM2000556	0.000	0.000			0.000	42.730	0.000 24.240	28.218
	NT2RM2000565	52.454	32.231			7.665	32. 508	18, 105	35.032
	NT2RM2000566	31.997				15.606	44. 727	22.394	28.766
	NT2RM2000567 NT2RM2000569	57, 110 113, 652	91.632				58. 576	40. 151	50.117
	NT2RM2000577	61.308	16, 114				83.608	36.221	60.695
E E	MT2RM2000581	152.797	45. 271				79. 582	62. 192	40.676
55									

Table 66

	INTARNACOCCA I	AC 161 T	03 700	104 000	37, 893	45.777	67 766	50. 428	50, 631
	NT2RM2000582	96. 163	83.789	104. 868			67.766		
	NT2RM2000588	109.847	89, 480	119. 521	70. 534	32. 168	143.491	88. 984	95. 908
	NT2RM2000589	91.130	45. 398	66. 143	21.774	22. 548	80.656	43.854	35. 379
5	NT2RM2000594	31.068	22, 138	28.684	10.809	13. 325	34, 179	10.310	16.391
	NT2RM2000599	275. 423	132.063	221. 911	86.738	66.363	237. 294	209. 381	119.304
	NT2RM2000609	26, 687	13.378	20. 025	9.729	14. 321	19. 395	17.956	8.545
	NT2RM2000612	40, 704	19.012	36. 338	9, 471	15. 531	27.049	24.872	30.269
	NT2RM2000622	45, 492	46.307	46.012	27.097	17, 426	48, 495	30.090	42.927
		279. 041	219. 374	245. 200	90.410	123.723	286. 194	221.925	144. 950
10	NT2RM2000623				60, 273	35. 044	29. 084	27.783	54.409
	NT2RM2000624	52. 551	88.174	87. 865					
	NT2RM2000632	15. 461	13.573	11.853	13.378	8.044	7.114	6.910	5.808
	NT2RM200D635	24. 725	21.442	42. 243	17.900	14. 353	23. 119	10.306	20.675
	NT2RM2000636	45. 247	47.662	62.828	24.460	33, 311	28.868	35.751	35.343
	NT2RM2000639	34. 707	19.290	26.594	15. 919	12.875	28.297	20.526	11.317
	NT2RM2000549	39.662	37. 102	62.088	31. 152	32. 252	42.335	27.796	50.424
15	NT2RM2000658	53, 598	26.723	55. 360	19.176	26. 348	46.815	23.949	20.812
	NT2RM2000660	84. 441	62.193	66.364	13.329	36.417	48.267	23.694	40.215
	NT2RM2000669	17.352	23.877	38. 180	11.181	16.885	17.594	13.008	20.479
	NT2RM2000689	118.126	102.565,	102. 237	102.435	37.057	156.147	96.539	140.413
	NT2RM2000691	29. 467	12.787	29, 631	9.783	15. 294	28.392	15.401	17.161
				122. 264	38. 290	60. 785	222.914	188.827	77, 434
20	NT2RN2000714	238.396	61.067					7.572	22.010
	NT 2RM2000718	9.515	10. 199	19.686	5.036	7. 922	8. 962		
	NT2RM2000732	44.022	24.869	42. 915	12. 209	29.863	38.537	30. 201	17.415
	NT2RM2000735	112.208	47. 966	111. 282	57.228	38. 980	78.590	45.888	59. 237
	NT2RM2000740	23. 990	62.438	143, 286	24.030	26. 159	35. 449	22.001	29.845
	NT2RM20D0743	15. 424	14.901	23, 591	12.391	9.779	16.339	8. 950	8.560
	NT2RM2000772	79.885	34.020	54. 908	31.068	31.256	64.893	44. 735	55. 557
25	NT2RM2000773	56.846	36.465	77. 155	26. 645	32. 523	60.130	42. 946	53.958
	NT2RM2000776	56.550	40, 820	69.793	43.736	22. 285	89. 348	33.285	45.221
	NT2RM2000784	54. 586	33, 888	45, 181	19.559	21.292	43.103	25, 540	42.124
	NT2RM2000795	169.462	132, 660	456. 283	117. 450	94. 702	91.566	59.832	91.914
	NT2RM2000796	12.942	12.033	20. 129	5.817	6.070	11.596	8, 538	11.009
	NT2RM2000798	67. 292	147. 984	71.980	42.802	43. 127	85. 427	63.126	132.706
30	NT2RM2000801	145.709	152. 451	160.966	85, 365	73,827	214. 221	157. 384	174.371
	NT2RM2000821	29.716	25. 994	36.976	14. 293	9. 638	63.476	12. 133	3.427
	NT2RM2000829		36. 834	148.015	32.077	69.569	70.012	26. 103	73.222
		77.695		51.022	19. 432	22.405	48.733	36.614	45.277
	NT2RM2000837	85. 748	27. 100					55.870	42.226
	NY2RM2000924	41.170	22.739	31.818	6.582	16.935	130. 595		
	NT2RM2000930	45, 514	31.120	39. 165	20.017	17. 433	49.111	28. 135	30.171
35	NT2RM2000937	85.092	19.912	28.613	13.728	34. 425	55. 176	53.959	15. 755
	NT2RM2000939	63.956	41.986	59. 137	18.909	23.056	57.088	26.370	29.465
	NY2RM2000942	141. 275	345.015	119.378	242.434	78. 282	274. 472	112.054	435.171
	NT2RM2000951	32. 383	20.717	32.763	17.041	10.179	32.704	19. 494	30.498
	NT2RM2000952	33.16D	18.882	34.052	15.194	27.783	44. 540	16.881	31.012
	NT2RM2000966	54.007	44. 546	57, 551	30. 397	27. 965	78.353	44. 947	77.916
40	NT2RM2000973	96.188	97.082	100.373	31.654	38. 259	115. 479	60.146	151.200
	NT2RM2000983	66.024	27.357	40.970	16.277	25.768	44. 322	40.901	34.882
	NT2RM2000984	38.635	39, 635	42.628	14.734	10.729	39.002	24.661	39.000
	NT2RM2000994	38.406	43.907	36.416	29, 495	24.408	22.384	18.679	31.517
	NT2RM2001004	74, 509	45, 438	146,622	36.919	35.918	125, 242	81, 529	92.360
	NT2RM2001022	195.677	346.056	350.501	243. 410	179.341	419.711	214, 981	540,668
45	NT2RM2001035	23. 201	26, 826	34.867	15. 930	11.692	19, 371	11.576	23.987
45	NT2RM2001038	18.846	16, 860	28.577	14. 251	9. 432	21, 182	12.726	12.544
	NT2RM2001038		17. 293	22.001	11.462	11.232	18. 219	25. 898	31.106
		31.149						49. 621	79. 938
	NT2RM2001050	101.638	45.617	56.097	28. 126	32.674	61.600		
•	NT2RM2001055	83.075	29.856	49.927	15.739	32.251	60.461	35. 925	29. 242
	MT2RM2001065	21.456	21.970	40, 162	20.006	27.398	26. 370	15. 034	14. 433
50	NT2RM2001075	366.658	258. 334	337.690	128. 945	166. 931	370, 161	257.064	228. 430
	NT2RM2001083	230. 683	79. 913	107.950	30. 576	63.142	203.365	79. 590	24. 253
	NT2RM2001100	182.772	114. 627	137. 289	65.878	54.062	141.899	155, 507	119.434
	NT2RM2001105	101.949	70.116	95.624	50.863	39.812	104. 272	87.573	85. 122
	NT2RM2001109	48. 591	27. 328	30.825	11.569	12.495	53.494	34.958	45. 222
	NT2RM2001110	99.871	68.967	152.982	31.616	42.715	78.028	71.894	63.509
55	NT2RM2001126	57.602	33.922	47. 638	18.667	20.095	52.257	42.378	28. 204

Table 67

	NT2RM2001131	59.454	21.547	32.934	24.063	22.706	37.676	28.873	17.418
	NT2RM2001141	116.250	82.599	275, 090	51.756	53. 514	85.069	47.274	63.199
	NT2RM2001152	20.261	21.814	23. 297	10. 506	9, 194	20.068	10.068	22,007
5	NT2RM2001177	44.847	43.449	52. 307	26.604	19.552	41.709	26.283	55, 231
	NT2RM2001T94	164. 727	54. 905	97. 293	28. 358	44.057	146.597	99.019	118.605
	NT2RM2001195	36.939	36.245	34.818	15.750	15.727	32.602	21.861	34. 274
	NT2RM2001196	125. 134	23.362	52, 729	15. 781	25, 090	77.518	62.058	31.794
	NT2RM2001201	56. 981	42.504	62.447	20, 139	31.351	68.607	32.835	44, 422
	NT2RM2001221	65.764	32.746	40. 357	19. 556	25.529	40.240	33.849	36. 497
10	NT2RM2001238	34. 807	25.200	33.023	13.254	14.872	43.011	20.155	18. 493
	NT2RM2001243	50. 316	49.076	42. 361	34. 148	33. 121	68.021	35.734	60.810
	NT2RM2001244	39.082	47.756	54, 069	35. 242	30.728	59. 908	22,778	50. 393
	NT2RM2001247	138.825	184, 906	146. 554	65. 082	57.954	94. 133	78.544	136.745
	NT2RM2001256	28. 147	18.773	29, 336	14. 133	9.881	8.739	16.106	25. 473
	NT2RM2001269	21.655	19. 444	36.676	14. 235	17.978	11.919	14.441	17.847
15	NT2RM2001278	105. 133	67.683	225, 135	41.243	42.803	61, 361	51, 930	64, 103
	NT2RM2001291	21. 264	19.798	31. 162	8.619	11.535	15. 945	16. 243	12. 482
	NT2RM2001294	60.754	44.696	66, 102	25. 820	20.715	42.950	28, 321	33. 134
	NT2RM2001295	43. 856	35.189,	40.675	10.220	16.301	35. 694	20.908	35. 879
	NT2RM2001302	30.816	16.802	26.058	10. 228	12.245	25. 513	14, 404	12.416
20	NT2RM2001306	11.584	52, 176	16, 722	6.379	6.616	13.560	8.347	10.145
20	NT2RM2001312	33. 361	18.866	54. 572	11.148	10.119	13.848	8. 526	26.714
	NT2RM2001319	13. 127	22.841	- 23.586	17.119	10.492	18.998	4.495	36.587
	NT2RM2001324	103.673	83.091	165, 198	32.861	22.835	56.112	31.793	39. 459
	NT2RM2001345	49. 534	25.168	35, 284	14. 837	16, 900	100.618	25, 540	19. 919
	NT2RM2001360	74. 152	33.097	38. 122	17. 360	16.021	50.562	31.265	21. 915
25	NT2RM2001370	28.821	12.859	21.986	6. 327	5.734	26. 406	10.631	2.394
2.5	NT2RM2001391	16.127	5.412	27.834	4. 575	4, 553	14. 188	3.910	9. 994
	NT2RM2001393	57.930	25. 241	58.135	14. 781	20. 544	47. 187	32.903	28. 104
	NT2RM2001420	17. 272	10.676	16.079	6.774	6.751	2.717	3.157	8. 464
	NT2RM2001423	17. 345	9.837	15.261	12.233	6. 527	15. 432	10.007	10. 935
	NT2RM2001424	196.973	74.966	136.019	35. 222	48.814	142.268	95.111	56. 187
30	NT2RM2001482	265.035	123.493	274. 926	59.811 28.605	19.730	227. 572 68. 321	99. 155 23. 722	72. 372 26. 475
	NT2RM2001499 NT2RM2001504	65. 942	48.790 24.742	62.383 30.958	9. 395	16.991	46.880	13.034	16. 709
	NT2RM2001524	39. 282 24. 755	14.244	24. 384	9.699	10.204	16. 924	9, 647	14. 539
	NT2RM2001530	5. 573	8. 914	10.768	5.856	3. 286	9. 623	4. 337	7.511
	NT2RM2001533	69. 137	57.026	127, 055	29.970	34. 159	33. 371	27.483	25. 268
	NT2RM2001540	65.400	54.541	73.017	63.277	35.636	49, 097	31, 308	76.346
35	NT2RM2001544	18.067	19.624	25. 228	12.549	7.049	19.380	11.033	9. 485
	NT2RM2001547	22.357	25.608	19.122	11.755	13.130	14. 503	12.339	10.697
	NT2RM2001558	59.623	25.861	31.696	14.111	16.568	53.758	34, 506	18.325
	NT2RM2001575	53. 128	46. 425	111.368	27.392	24.257	43.005	25. 405	24. 423
	NT2RM2001582	59.050	42.778	132.294	24. 555	24.449	28. 347	22. 303	22. 397
	NT2RM2001588	35. 342	21.815	27. 343	8.806	14. 132	21. 498	16.451	22.464
40	NT2RM2001592	19. 456	18, 542	28. 435	10.182	12.538	15. 234	15. 478	15.460
	NT2RM2001603	42.456	15. 253	41.037	12. 377	16.738	23. 117	21.517	12. 277
	NT2RM2001605	60. 434	36.233	43. 204	13, 580	20.116	41.260	20, 117	15. 459
	NT2RM2001611	54.771	39.056	128. 984	17.180	24. 100	40.047	19.191	16. 136
	NT2RM2001613	39.500	22.894	27.579	12.321	45. 138	26.696 168.993	21.149 96.729	24. 773 42. 842
45	NT2RM2001626 NT2RM2001632	202. 358	40.774	93.458 47.586	19.731 25.780	18: 848	32. 974	21, 939	45. 513
45	NT2RM2001632	30. 160 5. 521	9.885	12.546	7.571	6.017	11, 226	7. 294	20, 798
	NT2RM2001635	188.515	41.783	101.462	30.227	41.863	115.049	88. 246	58. 313
	NT2RN2001636	26.880	23.087	31.788	15.679	14.225	22.589	16.870	26. 264
	NT2RM2001637	13.020	5. 524	6.631	4.897	11.170	10,700	20.526	5, 331
	NT2RM2001639	71.531	28.740	32.389	12.149	15.813	54.897	28. 931	13. 443
50	NT2RM2001641	39.297	32.462	49.334	14.630	22.002	30. 556	21.763	16.776
	NT2RM2001643	25. 535	12.621	15.764	6.658	12.027	21.274	22.136	12.847
	NT 2RM 200 1648	26.584	18.351	24.507	8.310	6.636	18. 218	14.277	13. 561
	NT2RM2001652	18.655	15, 854	22.304	6.782	9.644	25. 729	7.851	20.144
	NT2RM2001659	16.893	10.861	16.538	3.750	4.964	9.228	6.172	11.278
	NT2RM2001660	17.414	13.987	20.619	12.709	10.544	12.482	10.671	11.244
55	NT2RM2001664	32.470	29, 186	27.804	16.171	15.728	29.928	13.136	17.877
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Table 68

	NT2RW2001668	89.325	61.356	52, 221	32.055	34. 144	88.196	46.704	35.968
	NT2RM2001670		20. 552	40. 552	17, 717	15, 452	67.725	25. 514	50. 962
		58.448	15. 752	21.018	19, 530	8. 980	62.746	15. 913	35. 807
5	NT2RN2001671	31.368	7. 210	6. 726	2. 026	5. 059	4. 678	4, 675	6.219
	NT2RM2001675	7. 281		11. 234	3. 584	7. 454	5. 095	5. 298	21.547
	NT2RM2001681	6.784	7.472	22. 146	9. 525	9. 058	19, 334	21. 485	26.746
	NT2RM2001685	28.752	21.105	43, 734	11, 154	11.656	30.491	20. 238	33, 991
	NT2RM2001688	35.233		239. 543	46. 305	60. 347	69. 201	35. 903	61.706
	NT2RM2001695	82.068	103. 403		31.652	35.701	110.799	51.358	52.359
10	NT2RM2001696	101.355	65. 027	68. 822		31. 134	111.891	66.042	79. 232
	NT2RM2001698	146.791	45. 033	83. 099 25. 919	28, 886	8. 984	19, 132	14. 447	42.307
	NT2RM2001699	24. 737	20. 994		3.702	1.312	7.813	9. 485	4. 374
	NT2RM2001700	14.734	8. 383	12. 975 50. 059	14, 943	24. 129	36.190	27.006	62, 522
	NT2RM2001704	50.393	27.867	144. 702	48. 167	41.996	55.679	28.916	57.741
	NT2RM2001706	75.476	62.308 12.916	25. 654	7.345	10.946	13.341	5. 670	10.957
15	NT2RM2001714	14.876	99.615	122.970	48. 569	68.313	188, 154	109, 177	48.112
	NT2RM2001716	294.058		57.895	15.717	34.379	105. 548	62.864	26.050
	NT2RM2001718	109.052	48. 161 14. 923	16.575	7, 233	8. 937	39.809	8.807	9. 491
	NT2RM2001723	20.352	41.046	46. 272	22.665	16.545	51. 332	33. 590	46.539
	NT2RM2001727	57.044	22. 859	24.865	8. 552	9.397	19.553	13.897	12, 427
	NT2RN2001730	27.206 25.036	6. 229	18.054	7.967	10.452	22, 532	14, 238	26.610
20	NT2RM2001738 NT2RM2001743	31, 219	15. 575	27. 495	8. 999	13.856	19.966	21. 123	40.203
	NY2R#2001743	41.699	57. 379	66. 833	29. 155	36.474	48.608	37, 342	50. 583
	NT2RM2001755	102.308	95. 543	95. 880	48. 800	50.926	85.016	46. 946	58.535
	NT2RM2001760	36.852	29. 592	43.280	11.529	16.235	41.973	21.095	36.897
	NT2RM2001765	17.310	22. 525	20.809	5. 472	6, 161	36.420	11.083	21.129
	NT2RM2001767	507.383	198.624	244.752	82. 225	86.662	313.630	261.579	156.449
25	NT2RN2001768	14.334	16.852	22. 405	14.516	7.327	13.653	4. 371	27.736
	NT2RM2001771	33.884	31.815	59.888	20.959	19. 261	40.562	26.114	70.587
	NT2RM2001778	14.653	9. 177	12.741	0.999	6.577	9.552	8.651	6. 525
	NT2RM2001782	49.540	17.667	39. 944	11.809	19.235	60. 433	38. 302	42.078
	NT2RM2001784	31.529	23.807	34. 905	9.620	16.512	26.774	14.749	17.008
	NT2RM2001785	73.444	32.799	54.722	14.868	28.332	74.431	52.678	40.155
30	NY2RM2001792	82.550	48. 689	54.661	13.880	26. 470	67.309	56, 934	51.170
	NT2RM2001795	130.534	65. 803	79.887	22, 935	40.781	108.971	66.672	68.900
	NT2RM2001797	17.770	23.911	46.302	31.918	15.965	38.330	15. 267	61.440
	NT2RM2001800	32.076	15.750	32.039	9. 323	10.196	25.569	24. 848	32.579
	NT2RM2001803	18.883	19.806	27.862	15.915	15.790	17.317	12.178	25. 827
	NT2RM2001805	10.973	6.105	12.362	3.395	7.748	17. 242	7. 464 18. 192	10.576
35	NT2RM2001806	41.604	28.683	30.345	12.360	14.554	35. 269	7, 004	10.764
	NT2RM2001813	11.155	10.752	12. 187	5.926	6.671	17.463	12.571	9. 506
	NT2RM2001814	16.422	18. 276	19.059	5, 168	10.179	28.082	23. 903	16.747
	NT2RM2001818	37. 340	15. 047	25.378 12.712	7.050 4.562	7.791	10.847	8. 727	7.819
	NT2RM2001823	13.814	13.268 37.936	22.505	15. 145	17.486	21.050	17. 161	33.945
40	NT2RM2001825	27. 524 68. 657	29.677	30. 202	9.749	22. 522	37.241	30, 727	18. 205
40	NT2RM2001839	53,715	31.908	39. 273	13. 944	12.144	27.291	25, 952	18.815
	NT2RM2001840	108, 411	98. 429	259.021	48.048	32.857	58.314	28. 523	37.338
	NT2RM2001851	52. 202	39, 752	63.088	24, 308	18.778	32, 821	26. 626	85.668
	NT2RM2001855	33.026	24. 176	29.953	16.912	19. 394	23.562	31, 355	25.910
	NT2RM2001867	30.838	22. 957	35. 457	14.948	16.183	32.799	17. 562	46.800
45	NT2RM2001869	129. 599	162.083	180. 222	173.694	64.737	231.277	145, 176	147. 129
<b>4</b> 5	NT2RM2001879	14, 477	14.016	20.104	6. 241	7.997	18.463	6.634	17.934
	NT2RM2001883	42.649	14.914	59.041	11.657	28.809	14.670	17, 172	5. 396
	NT2RM2001886	31.621	19.917	31.650	19.861	14.683	19.396	24. 519	18.912
	NT2RM2001887	19.995	18.787	31.384		6.192	7.945	11.032	6.537
	NT2RM2001896	5201.332	1475.462	2605.875	738.729	3013.651	5911.225	5347.627	1306.593
50	NT2RM2001902	9.512	5176	9. 030		3, 539	7.418	7, 583	3.383
	NT2RM2001903	63.243	40.127	55, 162		22.732		28.595	48.438
	NT2RW2001930	108.255	64.649	109. 195		39.123			59.426
	NT2RM2001935	36.519	23.148	18.415		11.165			10.042
	NT2RN2001936	78. 536	47.939		18. 980	21.013		35. 554	52.419
	NT2RM2001939	23.961	5. 651	21. 192					5. 482
55	NT2RM2001941	71.450	49.630	78.923	19.738	22.274	54.128	31, 260	34.949

. Table 69

	NT2RM2001950	46.415	29.816	36.996	18. 559	8. 239	39. 347	13.956	23. 224
	NT2RM2001952	2.871	2.885	10. 523	6.195	0.000	2.538	1.846	8. 237
5	NT2RM2001976	42.702	29. 344	52.698	20.599	18. 125	57.645	24. 197	33.972
3	NT2RM2001982	20.947	25. 776	25. 162	18. 275	10.576	18. 050	9, 191	14.830
	NT2RM2001983	23.643	16.045	27.861	9.316	13.749	23.964	18. 258	17.035
	NT2RM2001984	147.043	51.662	81.658	22.066	35. 725	120. 259	90.102	44. 130
	NT2RM2001989	76.106	50.939	80.150	44. 331	24. 785	39.074	34. 205	61. 176
	NT2RM2001996	37.798	41.931	43. 246	23.083	19.109	45.858 58.320	26.665 33.747	31.923
10	NT2RM2001997	63, 158	41.928	28. 543	20.691	22.046	45. 674	22.546	20.062
. •	NT2RM2001998	47.869	29. 374	50.969	17.042	23. 450	26.601	19.613	27.167
	NT2RM2001999	23.045	23. 925	34.107	16. 137	19. 923	57.270	34. 588	36. 304
	NT2RM2002003	60.554	45. 534	133.518	30. 271 8. 483	9. 918	13.788	12. 592	4. 939
	NT2RM2002004	16.782	14.896	24. 193 37. 573	16, 205	17. 990	24.047	10.371	17. 159
	NT2RW2002009	22.784	26.292	20.605	9.676	8. 686	10. 127	8.085	17.091
15	NT2RM2002014 NT2RM2002019	12.027 45.009	11.499	61.370	29.641	24.044	45. 990	20. 852	29.924
	NT2RW2002019	100.329	58. 955	73.738	25.096	36.513	90.878	44.848	41.854
	NT2RM2002023	53.030	36. 122	48.637	23. 542	18. 217	49.856	26.265	32.557
	NT2RM2002034	55.319	58.655	69.310	15,775	34. 969	119.355	37.851	31.637
	NT2RM2002049	30.306	26. 333	67.224	12.461	13.486	32.196	19.763	26. 143
	NT2RM2002055	4.746	9. 322	10.601	1.587	3.475	2.738	4.711	1.253
20	NT2RM2002072	274.106	142.825	221,668	99.170	111.051	240.179	193. 919	147.089
	NT2RM2002088	66.101	43.548	67.009	20.108	29.769	38.434	34. 203	37.710
	NT2RM2002091	157.752	95. 255	103.301	42.530	51. 107	91.971	64. 745	58. 827
	NT2RM2002100	36.481	42.661	83.563	34, 382	22.604	42.960	30, 266	45. 203
	NT2RM2002109	65.961	25.178	54.629	11.426	17.601	56.066	33. 542 195. 547	37. 167 168. 748
25	NT2RM2002126	271.768	145.370	244. 199	79. 521	110.685	272. 182 20. 446	37.022	29.430
23	NT2RM2002128	30.978	20.989	35.773	13.699 14.507	15. 221 24. 022	54, 089	40. 427	15. 416
	NT2RM2002129	53.911	38.709	50. 544 127. 900	44. 871	54. 994	121.896	116.748	122.762
	NT2RV2002142	157.794 39.141	95. 271 23. 769	42.061	18. 362	18. 425	90. 424	32.619	22.086
	NT2RM2002144 NT2RM2002145	69.465	33.538	54.629	19.065	28. 804	64.861	31.013	26.312
	NT2RW2002143	57. 982	34.658	45. 808	37.204	22.363	85.615	33.858	45.468
30	NT2RM2002163	46. 164	22.611	32.853	10.533	12.313	28.767	18. 529	24. 578
	NT2RM2002170	20.367	15. 918	26. 954	17.854	7.659	21.614	6. 584	31, 812
	HT2RH2002178	72.826	29.934	35, 113	17.819	17.814	57.676	53.788	36.064
	NT2RN2002179	20.487	16.890	26.778	4.596	7.536	27. 483	11.691	22.514
	MT2RM2002270	75.965	30, 835	59.481	19.162	23. 264	67.579	38.824	31.179
	NT2RM2002326	25.054	17.109	25.901	10.631	13. 295	20.170	15. 155	11.219 36.266
35	NT2RM2002337	49.608	30, 430	44. 382	14.424	20.214	49. 783 129. 046	38.536 67.853	46.026
	NT2RM2002339	126.783	46. 855	62.446	22.680 17.636	35. 280 9. 930	27, 503	20.940	24. 302
	NT2RM2002345	34.662	27. 251 67. 271	30. 489 118. 627	55. 152	36, 416	61.876	35. 957	79.909
	NT2RM2002368	53.018 29.049	17.380	20. 958	5. 965	9.584	35.715	13.371	27.731
	NT2RM2002424	23.738	30.901	58.344	39. 153	17.434	49.766	25. 216	77.325
40	NT2RM2002450	40.370	29.535	54. 082	14. 242	16.219	34. 988	19.676	33.464
40	NT2RM2002482	44.705	26.737	46.955	14.769	18.437	42.664	46.045	30.188
	NY2RM2002492	113, 197	127.579	109.738	72.932	49, 321	103, 335	74.905	97.173
	NT2RM2002575	112.457	88.605	247.074	59. 323	48. 212	80.685	45.794	62.455
	NT2RM2002580	64.838	62.853	111.962	57.513	25. 109	65. 998	30.240	96.173
	NT2RM2002592	110.441	70.152	96. 103	45.340	44. 856	104.438	69.434 17.823	61.212
. 45	NY2RN2002608	20,462	46.581	29. 949	14. 231	13.430	45. 176	71.069	33.280
	NT2RM2002615	33.564	24.375	25. 868	12.468		108, 504		91.258
	NT2RM2002622	95.365	53.669	62.071 276.792	68.615	58.079		51.946	80.285
	NT2RM2002630	118.784	86.444 30.749	31.925		20.353		32.736	22.117
	NT2RM2002634 NT2RM2002645	36.887 51.215	209.069	58. 292			97.660	24.132	61.537
	NT2RM2002646	69.318	57.452	61.629		19.295			24. 267
50	NT2RM2002647	31.140	27.535						42.269
	NT2RM2002652	42.576	30, 866	34. 782		12.829	46. 172		30. 578
	NT2RM2002692	53.871	40.724					30.444	71.284
	NT2RM2002721	81.740	78. 721	123. 105	75. 203			44.593	72.005
	NT2RM2002748	91.982	206.064						228.776
55	NT2RM2002764	46.071	41.769	48. 814	22. 081	22.119	34, 365	32.761	36.777
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Table 70

	CUVACUASAS TES	AA AAF 1	40 044 1	68.101	23.056	28, 389	72.818	41.505	60, 302
	NT2RM2002772	80.296	40. 944						
	NT2RM2002811	63.439	38. 909	43.044	17.983	20. 375	56.523	23. 815	28. 434
	NT2RM2002818	50.605	52.430	151.915	32, 193	19.702	26.680	17. 380	40.512
5	NT2RM2002879	24.562	28.586	34. 172	8.860	5. 095	18.514	12. 159	30.354
	NT2RM2002979	84.387	41.192	53.776	21.436	31.083	74.067	53.736	47.429
				33. 191	11.478	15. 597	54. 899	35. 830	32.861
	NT2RM2002981	59. 340	25.706					30, 448	42.538
	NT2RM2002995	42.179	21, 303	31.267	13. 206	10.830	32.109		
	NT2RM2001031	44.114	29.430	46.063	16.774	17. 437	43. 222	40. 155	25.053
	NT2RM2003042	106, 509	160, 917	155. 488	83.058	73. 174	152.473	69.308	122.583
10	NT2RM2003044	33. 909	33.603	47.142	12.698	45, 517	25. 310	25. 508	29.529
		47.953	25. 520	41.051	9.604	15. 180	34, 197	23. 552	25,659
	NT2RM2003090				11.759	18. 398	29. 592	34. 566	28.874
	NT2RM2003095	43.943	31.580	32.103				12. 203	
	NT2RM2003116	20.590	18. 126	22.701	10.734	10. 194	11.727		14.479
	NT2RM2003222	21.398	10.313	27.148	5. 349	13. 395	13.068	20. 550	25.145
	NT2RM2003224	110, 266	37, 406	48.819	30.835	29. 947	80. 454	57.677	53, 588
15	NT2RM2003250	30.062	26.498	38.776	15,773	16.547	23. 997	24.660	26.915
		12.707	12.077	15.752	5. 247	7.979	8. 239	5.752	8.852
	NT2RM2003258					33. 378	31. 965	36.375	43.803
	NT2RM2003262	37. 575	42.567	50.603	27.374				
	NT2RM4000023	49.690	44.882	57.421	17. 352	24.868	53.007	25.083	35.943
	NT2RM4000024	33.710	23.142	26.564	7.803	10.308	34. 975	25. 466	17.156
	NT2RM4000027	6.576	5, 402	9.541	2.488	3.969	5.783	1.681	9.230
20	NY2RM4000030	107. 340	43.649	64.579	25. 595	27.984	81.398	45.801	45.851
•	NT2RM4000033	54. 521	41. 188	116.087	19.883	18. 324	28.028	14.764	29.244
			20. 135	21.495	9. 212	9. 086	13. 100	7. 920	12, 176
	NT2RM4000034	8.646						23. 736	18.823
	NT2RN4000046	42.055	17, 446	23.148	8. 687	9. 540	32. 532		
	NY2RM4000052	23.740	17, 236	25.146	8.065	5, 341	17.707	13.080	13.561
	NT2RM4000054	440. 502	221, 475	352. 543	107. 153	132.322	410, 274	281.112	209.475
25	NT2RM4000061	30.264	15.792	27.807	6.396	10. 845	21.557	14.902	4. 276
	NT2RM4000074	8.073	35. 126	41.073	20.510	9.480	34. 431	24. 493	47.368
	NT2RM4000085	22. 897	19.315	23.277	16.541	12.977	24, 111	12.451	24,618
					20. 299	18.217	28.085	16.663	27.361
	NT2RM4000086	50.715	22.670	78.725					12.129
	NT2RM4000100	17.872	21.935	15.019	10.707	10.091	15, 556	12. 260	
	NT2RM4000101	42.770	15. 130	25.674	6. 552	7.785	24. 576	15. 561	5.064
30	NT2RM4000102	407.848	190.329	321.537	152.733	208.613	334, 316	212.009	231, 229
00	NT2RM4000104	23.885	13.626	17.310	3.131	7.950	21.156	10.845	7.969
	NT2RN4000115	32.088	10.072	16.134	5. 693	9, 226	13, 512	10.582	7.588
	NT2RM4000129	36.681	21.490	22.965	12. 521	11.849	23.308	16.146	10.761
			23.620	31.564	24. 507	22.610	18, 556	14.008	44.620
	NT2RM4000139	25. 930							42.075
	NT2RM4000149	33. 404	17.925	29.734	13.712	15. 989	18. 474	26.736	
35	NT2RM4000155	21.566	44.820	46.750	15. 598	16.524	14. 928	9.733	8.224
	NT2RM4000156	16.586	6. 239	5.822	3. 387	3. 958	28.594	7. 207	15.119
	NT2RM4000167	20.171	16.879	15.859	11.667	2. 739	8. 443	3.474	21.050
	NT2RM4000169	30.428	28.089	36.443	24. 244	11.338	20.566	13.227	60.152
	NT2RM4000191	52.656	25. 321	40.946	12.980	18.787	41.092	35.047	38.394
	NT2RM4000197	15. 240	11.946	16.612	2. 282	13. 434	15. 387	8. 823	5.757
							49. 371	53. 195	32,774
40	NT2RM4000198	88. 525	63.904	196.728	39. 099	37. 803			
	NT2RM4000199	52, 380	24.904	46.280	17. 110	18. 960	33. 287	27. 322	30.945
	NT2RM4000200	33. 395	16.462	28.537	10.600	16.103	20.714	14.030	6.949
	NT2RM4000202	30. 208	20.922	42.468	9. 182	9. 970	16.908	10.274	12.811
	NT2RM4000210	66. 407	27.815	30.474	15. 335	15.812	41.212	27. 389	47.172
	NT2RM4000215	25.869	24.845	36.251	22.848	13. 152	31.488	12.403	27.548
	NT2RM4000220	47. 201	39.573	38.877	20. 267	19.583	51.592	35.424	51.912
45						14, 171	29. 316	28.242	16.590
· <del>-</del>	NT2RM4000229	38.395	26.396	42.302	13.878			28.746	34.406
	NT2RM4000231	54.697	33, 959	43.440					
	NT2RM4000233	209. 479	90.187	137.270	36. 159	66.994	160.853	100.732	62.965
	NT2RM4000244	16.916	9.010	13.401	4. 357	9. 911	12. 907	8.771	8.963
	NT2RM4000251	43.833	19.474	33, 500		16.673	31.966	32.833	8.105
	NT2RM4000255	35, 799	17.398	36,446		12.098	29.741	23.847	15.929
50			79.778	222, 138		51.026	72, 136	39.083	49.420
	NT2RM4000265	102.046						162, 352	166, 824
	NT2RM4000283	285. 571	172.391	189.067		94. 953	255. 306		
	NT2RM4000284	23.615	36.279	30.562		17.835	25. 501	27. 248	34.927
	NT2RM4000290	74.673	36.513	57.081	15.623	22.008	73.912	45.709	43.178
	NT2RM4000295	24.000	18.871	22.693	8. 987	11.022	47.890	18.701	14.976
	NT2RM4000306	140.029	42.148			78. 561	140.760	92.030	34.220
55		1 ,30.025	1 140	1 -1.411	1 . 3 30				·
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Table 71

	NT2RM4000307	20.678	19. 168	22.141	9.050	9.145	23.385	14. 143	13.754
	NT2RM4000309	41.662	20. 618	26.408	8. 581	10.787	30.894	18.116	11.868
						12. 239	39.520	34.145	43.040
	NT2RM4000313	36. 434	20. 403	33. 260	17.080				
5	NT2RM4000318	52.262	31.467	139. 471	20.774	17.880	23.820	17.441	19.608
	HT2RH4000324	51.333	27.748	39.958	9. 932	17.995	63.248	27.625	42.800
	NT2RM4000326	32, 179	16.471	20.536	8. 435	10.621	23.791	17.926	20.620
	NT2RM4000327	60.230	58. 958	198.666	39. 302	28. 376	44,008	20.961	43.734
	NT2RM4000344	63.708	65.489	173.360	18. 949	27.536	34.270	15, 519	42.106
	NT2RM4000349	30.022	14.663	14.070	7.442	10.197	22.535	12.455	16.210
10	NT2RM4000354	46.698	15. 085	27.013	11.329	7. 922	27.895	13.694	15.005
	NT2RM4000356	32.497	24. 336	32.372	13.972	11.464	43.673	31.608	29.630
	HT2RM4000366	528. 262	330.865	423.109	167.985	170.232	378.411	215.606	442.307
	NT2RM4000368	51.220	51, 300	153. 236	33.445	22.538	43.253	17.539	64.383
	NT2RM4000373	25. 297	22.861	32.020	19.516	16.128	25.045	13.784	37.614
	NT2RM4000385	22.576	9. 738	24.078	8. 987	9.704	21.730	24.414	23.758
15	NT2RM4000395	61.364	79, 696	124.563	37.133	40.433	107.248	46.227	46.047
	NT2RM4000414	159.474	59, 130	69.911	18.566	40. 333	119.002	79.051	21.561
	NT2RM4000417	15.712	20.634	23. 502	7.213	7.502	15.030	7.412	1.857
	NT2RM4000421	15.106	14.708	19.062	8.549	6.469	15.114	8.074	20.588
	NT2RM4G00425	101, 441	83.854	259. 486	55. 511	39, 319	53.250	31.739	69.026
	NT2RM4000423	51.457	24. 650	39.654	12.379	16.608	41.753	37.139	36.708
20	NT2RM4000436	51.207	21.755	29. 307	13.444	12.333	34.290	27.223	37.320
	NT2RM4000444	40.864	26. 268	67.826	11.797	17,600	39.060	23.113	28.672
	NT 2RM4000457	63.983	39.080	61.124	23. 292	28.748	50.040	28.813	31.965
	NT2RM4000473	41.652	29.088	37.803	8, 939	15.093	35.469	20.877	14.796
	NT2RM4000472	68. 502	62.226	206.357	48. 752	23.646	77.597	28. 412	104.099
•	NT2RM4000486	30.140	26, 427	28. 452	18.097	7.542	22.184	12.697	24.533
25	NT2RM4000490	51.124	23. 641	42. 235	9. 300	14.683	56.785	25.625	17.105
23	NT2RM4000496	110.770	31.642	65.060	13.739	27.500	68.720	52.247	37.631
	NT2RM4000505	134. 100	84.063	126.035	43.665	56.053	130.720	81, 120	71.520
	NT2RM4000511	73.441	160.671	81.146	172.018	35.906	98.128	55.037	164, 299
	NT2RM4000514	24.804	23.670	34.085	13.945	16,589	32.103	21.758	11, 170
	NT2RM4000515	56.528	99, 798	88.516	40.030	41.279	67.061	40.210	72.202
	NT2RM4000517	94.295	97.384	143. 107	76. 451	43.905	144.940	69.520	145.604
30	NT2RM4000520	13.459	13.780	16. 902	5. 273	5.564	7.899	7.054	14. 968
	NT2RM4000531	29.188	24. 283	26.738	11.063	12.826	18.929	23.443	20.712
	NT2RM4000532	14. 395	12.711	19. 277	9. 437	8.520	12.914	15.215	13.835
	NT2RM4000533	18.380	13.704	18, 165	8. 534	7.454	15.515	10.288	7.686
	NT2RM4000534	17.803	11.768	18.975	7. 585	10.236	14.119	11.420	19.497
	NT2RM4000563	53.983	34.056	51.401	17. 700	36.352	45,609	32.373	33.367
35	NT2RM4000566	36.586	22. 989	35.859	9. 957	21.078	25.668	24. 949	21. 224
	NT2RM4000568	59.423	29.845	36.652	12. 139	25.850	70.617	54.001	29.192
	NT 2RM 4000585	48.810	27.673	38.443	12.701	20.510	33.948	23.868	27.346
	NT2RM4000587	29.705	26.644	25.876	12.729	11.927	16.240	17.926	19.718
	NT2RM4000590	32.164	21.289	29.186	8. 941	11.617	18.856	16.495	13.544
	NT2RM4000593	61.080	32.766	38.970	15.411	20.350	33.032	30.484	25.715
40	NT28M4000595	41.141	22.473	35, 313	9.768	11.448	11.237	20.012	12.069
	NT2RM4000603	78.976	52.410	58, 176	24.839	24.042	50.072	40.363	31.910
	NT2RM4000611	15.953	10.734	13.469	9, 013	8.977	10.161	7, 157	22.979
	MT2RM4000816	45.814	37. 309	35, 175	17. 505	23.768	40.117	27.918	39.007
	NT2RM4000621	57.493	77.709	73.014	76.819	24.081	71.204	46.769	83.169
	NT2RM4000648	28.637	18.518	26.908	8. 210	13.083	15.965	12.644	11.022
45	NT2RM4000649	85.058	41.743	59.668	13.629	29.612	55. 983	39.586	36.405
	NT2RM4000658	135.688	61.028	120.722	28. 197	43.765	79.777	46.011	96.630
	NT2RM4000661	71.864	99.345	52.294	18. 409	29.132	62.897	45.030	41.904
	NT2RM4000673	135.680	61.584	75.017		20.618	70.048	46.608	45. 107
	NT2RM4000674	75.722	36.633	51.480			34, 561	42.749	30.664
	NT2RM4000689	41.790	28.540	39.966		8.448	22. 615	15.641	20.045
50	NT2RM4000698	61.169	46.347			41.257	63.885	38. 390	29.637
	NT2RM4000700	27.239	106, 106	27.114					12.082
	NT2RM4000701	227.264	115.040						65.330
	NT2RM4000712	43.183	27, 951						16.644
	NT2RM4000717	34.385	22.333						20.540
	NT2RM4000733	75.958	43. 996					65. 331	37, 193
55	111 TURNOUT 33	1 (3.330	1 -10. 530	45.560					

Table 72

			40 670 1	E3 736 I	12 070 1	12 156	20 047 1	23.333	39,227
	NT2RM4000734	24. 197	38. 270	53. 725	16. 970	13. 155	39.087		26.083
	NT2RW4000741	43.844	13. 589	30. 427	10.346	8.744	26.119	12.592	83.553
5	NT2RM4000744	50.833	14. 548	25.024	23. 480	10.805	62.136		100.669
3	NT2RM4000749	80. 902	71.083	91.533	27. 354	60.031	198, 030	52. 328	
	NT2RM4000751	22.688	29.768	53. 788	53. 315	27. 282	19.811	22.272	42.714
	NT2RM4000752	52.247	32.866	40.812	14. 427	15. 224	9. 355	23.407	43.927
	NT2RM4000760	33. 235	16. 169	27. 997	11.989	19. 412	13. 254	10.563	10.820
	NT2RM4000761	2403. 264	848.134	3887. 956				2359.029	400.128
	NT2RM4000764	301,709	144. 132	163.494	49.659	143.743	257.369	245.639	103.045
10	NT2RM4000768	11.747	9. 247	11.542	9.135	9. 038	10. 345	6.336	11.267
	NT2RM4000778	6.893	5. 725	9, 950	5.466	4. 458	5. 886	5.079	5.685
	NT2RM4000779	238, 073	96.516	182, 851	51.850	99, 170	184. 671	138.565	75.926
	NT2RM4000787	69, 121	57.977	157, 708	28. 426	29, 213	21.609	22.633	11.420
	NT2RM4000790	50. 309	46.026	83. 182	23. 988	30. 494	22.815	35.485	31.417
	NT2RN4000795	453, 425	108.548	204.710	17.809	92. 365	272.802	147.653	47.088
15	NT2RM4000796	144, 288	57.098	70.720	23.213	47.104	97. 550	50, 426	30.942
	NT2RM4000798	59. 938	28. 301	25.839	10.244	18, 327	23.444	20.572	11.548
	NT2RM4000800	150.768	122.487	195.880	137.376	57.284	146, 130	97.369	185.386
	NT2RM4000813	37.084	20. 876	36.294	12.655	14. 527	25. 975	22.848	11.921
	NT2RM4000820	86.855	60. 381	192.196	39.751	37.738	50. 427	35.797	26.747
	NT2RM4000827	41.788	28.006	51, 622	20.945	21.631	21, 541	30.438	31.570
20	NY2RM4000830	68. 078	30.965	59.647	20. 203	26.347	37.484	30.029	44.495
	NT2RM4000833	111, 407	74. 480	77.732	17.832	39. 802	56.697	25, 292	36.404
	NT2RM4000841	49, 942	45, 599	72.313	16.308	20.094	29.644	26.188	28.854
	NT2RM4000845	104. 561	76. 278	275. 932	57.490	49.037	63.058	35.772	14. 948
		125. 196	36.830	101.007	17.584	32.806	82.740	51.262	19.922
	NT2RM4000848	113.009	77.800	126. 539	43, 464	43.880	57.479	52.365	44. 156
25	NT2RM4000852 NT2RM4000855	64.608	50. 229	146. 326	22.844	23.661	28. 928	25.813	51.332
2.5	NT2RM4000859	24.418	19.759	24. 141	10.385	14, 916	34, 345	18,598	11.625
	NT2RM4000868	16, 564	14. 752	14. 556	11.565	9, 114	12.226	17.324	12.029
	NT2RM4000870	55. 531	47.020	57.796	18.791	30. 154	39.778	25. 127	26.057
	NT2RM4000879	103.887	41.773	56.495	12.837	31.154	67.942	43, 586	22.044
	NT2RM4000882	81. 982	42.561	80. 304	22.840	38.713	36.853	45.646	48.992
20		151, 731	36.758	112.092	22,645	40.960	98.527	85.229	22.008
30	NT2RM4000887	84. 679	41.293	172.935	28.755	27.724	44, 297	19.644	26, 291
	NT2RM4000897	45, 994	42.630	58. 329	17.578	25. 299	44.317	41.019	30.575
	NT2RM4000901	13. 138	13.528	18.046	7.930	5. 669	7.738	9.304	5.798
	NT2RM4000950	13.710	21.028	17. 402	10.585	11.390	13.090	8. 272	13.397
	NT2RM4000965	54, 459	35.282	50. 127	15, 952	25. 327	23.064	21.414	26.049
	NT2RM4000971	41, 258	27.847	39.604	12.433	17.061	72.230	20.025	17, 430
35		33.580	21.677	32.692	7.475	11. 547	22.259	16.549	12.389
	NT2RM4000979	51.537	23. 981	27.883	11.309	12. 974	42.714	19.808	18.064
	NT2RM4000989	43. 246	16.680	33.780	10.504	10.430	22, 581	33.282	15. 269
	NT2RM4000991	6. 595	8. 954	14. 910	4.216	4, 093	24. 193	3.472	15, 581
	NT2RM4000992	61.901	44.659	179.747	37.376	29. 327	33,667	22.750	38, 582
	NT2RM4000996	12.902	17.829	47, 104	22.304	9. 589	15. 133	12.379	41.017
40	NT2RM4000397	139.754	107. 958	216.478	45.750	59. 135	79.871	47.855	52.159
	NT2RM4001001	222. 229	90.117	123. 541	25. 902		102, 439	120.879	88.667
	NT2RM4001002	22.453	23. 223	34, 127	15.841	13. 942	17.616	10.393	26.669
	NT2RM4001016	39. 433	22.372	27. 844	7.677		29.791	22.346	14. 840
	NT2RM4001025	123, 159	184.713	262.665			167.042	104.628	258. 452
	NT2RM4001027	1,003	0.083	0.000			0.903	0.000	13.341
45	NT2RM4001032	15.446	8.560	20. 283	7.827		9, 129	9,798	10.321
	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	1 1 1 1 1 1	7. 922					9.323	18.916
	NT 2RM4001047	18,565	64. 640						27.816
	NT2RM4001049	87. 157 45. 597	65. 440						24. 356
	NT2RM4001051		54. 084					<del></del>	19.934
		83.704							
50	NT2RM4001053	55.548							14. 263
	NY2RM4001054	29. 223	12.533						64.703
	NT2RM4001059	181. 587	40.368						11.555
	NT2RM4001071	29.020	21. 136						
	NT2RM4001084	42.690							58. 081
	NT2RM4001092								46.701
55	NT2RM4001100	43.266	33.448	49.94	8.29	19.072	.   64.120	1 10. 221	1 70.101

Table 73

	MT2RM4001116	27.726	26.051	28. 521	6.793	9.001	18.038	14. 406	8.177
	NT2RM4001119	56.668	21.890	35. 980	9.796	15.859	38.916	35.588	15.608
	NT2RM4001140	136. 817	79.720	322.522	72.609	64.281	53.073	51.451	56.047
5	NT2RM4001148	238. 824	52.972	84.009	16.224	62.535	137.805	147.073	38. 797
	NT2RM4001151	49.119	18.810	31.963	9.013	16.522	24. 362	37.118	17. 496
	NT2RM4001155	51. 322	26. 524	38.663	9.832	19.192	16.401	24. 191	12.958
	NT2RM4001157	29, 926	19. 538	29.560	8.442	11.794	23.764	9.393	5.071
	NT2RM4001160	72.399	50.574	60.230	13. 285	29. 392	49.862	35.181	33. 807
	NT2RM4001163	150.688	70. 942	95.070	47. 204	58. 092	77. 447	65. 645	40. 117
10	NT2RM4001187	46.613	33.666	37. 323	10.669	19.756	22.493	19.909	13.410
	NT2RM4001191	62.821	78.568	138. 398	23.085	37. 250	19.851	28.068	31.505
	NT2RM4001200	48. 487	41.856	115. 958	43.120	35. 674	29. 433	29.755	46.933
	NT2RM4001203	29.740	33. 257	26. 183	10.711	18.414	17. 515	13.820	29.510
	NT2RM4001204	85. 368	2.729	5. 406	1.939	1.539	2. 503	5.782	1.987
	NT2RM4001217	22. 326	14, 483	20. 894	6.910	10. 252	17.142	14, 178	16.377
15	NT2RM4001245	102.964	61.341	59. 224	17.873	32.330	47.902	39.713	28. 855
	NT2RM4001247	60.472	48.248	105, 685	27.869	20. 131	20.633	22.912	17. 998
	NT2RM4001256	38. 132	20.867	27.791	11.662	11.297	22.362	18. 443	14. 221
	NT2RM4001258	13. 173	14.508	15.622	2.115	6.064	10.903	7.779	31.184 13.809
	NT2RM4001267	18. 994	10.887	19. 555	6. 271	8. 494	3. 421 21. 978	30.691	39.740
20	NT2RM4001273	57. 388	34. 293	59.413	25. 522 13. 708	17.714	31.923	19.522	23. 080
20	NT2RM4001281	52.686	24. 825 1240. 433	33. 241 782. 259	477.895	296.841	681.688	413.930	936.577
	NT2RM4001286 NT2RM4001290	481. 183 25. 298	23.154	13.373	6. 552	0.000	12.469	8.723	14.611
	NT2RM4001309	48. 445	24. 031	36.511	15.060	18.354	33.040	18, 409	21.487
	N12RM4001303	61.618	55. 950	171.030	27.704	18.541	31. 137	15.527	37.397
	NT2RM4001316	49, 175	40.348	93. 903	19.571	16, 907	28.903	20. 127	14.212
25	NT2RM4001320	73. 145	43.895	149.769	28.755	24. 031	24. 203	22.793	27.654
20	NT2RM4001321	49. 367	26.564	28. 912	10.370	15. 275	21.145	21.285	20.579
	NT2RM4001325	38.855	43. 433	53. 158	15.234	25. 333	31.524	26. 184	15.840
	NT2RM4001333	48.466	17.343	99.002	20.144	115, 167	148, 955	12.312	8.170
	NT2RM4001340	30.804	28. 992	40.576	27.062	32.009	10.155	18. 551	26. 573
	NT2RM4001344	30.624	35.092	33. 290	12.667	12. 525	9.910	11.004	11.417
30	NT2RM4001347	14. 549	14, 691	20.853	11.657	13. 229	14. 366	8.959	54.748
	NT2RM4001357	58. 256	26. 925	40.009	14.812	13.213	104.908	348.697	7. 592
	NT2RM4001360	86.062	33.099	53. 959	12.261	27. 140	48.858	36.604	20.008
	NT2RM4001371	57.075	37.841	49.730	24. 239	25.868 36.169	54.098 52.589	8.910 30.583	31.242
	NT2RM4001377	101.216	75. 138	68, 626 56, 186	19.407 35.607	24.700	70.227	41.803	66.511
	NT2RM4001382	56.509 13.506	78. 201 11. 432	7.793	6. 199	7, 970	12.881	6.788	7, 108
35	NT2RM4001400	21.837	16. 958	21.913	10.795	7.913	16.255	9.524	12.188
	NT2RM4001409	28. 309	17.011	26.656	9.796	12.960	23.632	14.054	20.949
	NT2RM4001410	29.072	19.001	30.576	8.925	14.550	18.489	21.014	17.448
	NT2RM4001411	8. 505	7.030	30.358	2.388	3. 324	0.962	1.969	1.931
	NT2RM4001412	59.413	25. 935	59.821	15. 231	22.577	30. 927	24. 563	11.190
40	NT2RM4001414	64.093	13. 321	33.046	9.873	26. 265	24.538	20. 805	20.958
40	NT2RM4001436	33.680	29.671	20.088	7.331	12.620	14.939	11,468	14. 185
	NY2RM4001437	70.569	41.529	158, 116	28.707	19. 302	25.565	23.649	23.787
	NT2RN4001444	63.099	33.815	51.190	21.250	36.920	56.421	7.237	35. 180 9. 931
	NT2RM4001454	15. 293	16. 251	33. 213	14. 589	11.226	13.235	13.432	28.743
	NT2RM4001455	8. 636	7.947	12.910	5. 235	6.864	7.007	22.451	46. 563
45	NT2RM4001483	74.168	64.931	192. 825	13.377	33.854 14.505	15.628	23. 221	19.361
	NT2RM4001489 NT2RM4001495	27.884	28.159 117.396	36, 108 133, 602			91.833	54. 255	51. 382
	NT2RM4001499	260.493 68.936	37. 210	73.295	19. 265	26.638	41.151	25,000	25.754
	NT2RM4001515	11.646	7. 906	18. 332		7.167	15.640	6.612	8. 512
	NT2RM4001519	12.556	9. 937	20.664			10.138	7.966	8. 328
	NT2RM4001522	71.440	69. 438	164.718			32.755	19.774	38.742
50	NT2RM4001523	24.710	16.532	29.750			12.279	19.569	31.077
	NT2RM4001550	24.908	22.060	34. 537			20.143	15. 284	28.090
	NT2RM4001553	73.682	40, 371	52.795		21.686	46.848	27.034	27.166
	NT2RM4001554	53.585	30. D46	33. 134			26.877	18,771	20.649
	NT2RM4001557	19.423	19. 434	24. 184			21.486	7.653	15. 404
<i>E E</i>	NY2RM4001565	65.552	37.852	90.440	18.538	17.294	23.128	23.413	18.529
55									

Table 74

				07 457	28. 565 1	28.860	79.976 1	52, 286	9, 785
	NT2RM4001566	100.946	48.659			8, 597	0.901	6.611	1.304
	NT2RM4001569	7.010	5.598	41.076	3. 288		35. 411	21.050	31.905
_	NT2RM4001579	41.258	24.859	37. 584	7. 247	15. 119	22.015	19.971	25. 442
5	NT2RM4001582	36.827	21. 162		10.109	10. 956	47. 320	41.167	35,619
	NT2RM4001589	57.574	32.795		23.877	20. 226	17. 425	7.965	10.850
	NT2RW4001592	32.950	21.429	32.007	7. 221	14. 392	46.488	34, 751	25, 706
	NT2RM4001594	55.970	26.805	46.827	13.556	21.275	51. 457	41. 254	42, 293
	NT2RM4001597	113.189		189. 284	36.307	35, 658	11.032	9. 672	10, 297
	MT2RM4001605	16. 347	11.965	18.084	2.805	4. 141	182.319		191.832
10	NT2RM4001609			265. 155	76.761	120. 584 38. 114	56. 107	36.218	28. 535
	NT2RM4001610	89.090	32. 924	55.024	13. 942	11. 242	12.351	22. 333	13, 486
	NT2RM4001511	30.709	14. 204	28.060	6. 394	28.633	44, 101	23.934	50. 341
	NT2RM4001618	77.313	59. 231	178. 569	26.795	35, 073	39. 451	29.062	30, 213
	NT2RM4001522	42.484	50.813	37.378	16, 153	11. 389	25. 162	26.300	19.356
15	NT2RM4001524	55.088	35. 243	39. 342	16. 243	29.704	87.349	62, 707	32, 707
13	NT2RM4001625	165. 457	44. 283	55. 076	10. 721	9, 407	17. 262	17.006	17. 599
	NT2RM4001629	23. 424	34. 729	31.319	80. 539	33.853	62.834	39. 339	102.299
	NT2RM4001632	49. 318	105, 740	25. 229	7. 187	11.536	12.746	15.743	15.315
	NT2RM4001642	25.758	83.479	257. 397	53.466	49.798	64.749	33.054	65. 546
	NT2RM4001647	140.643	17.016	26, 536	7.633	8, 417	10.663	14. 969	25. 969
20	NT2RM4001650	93, 433	61.261	62. 868	18.713	28.801	43. 545	39. 576	18. 233
20	NT2RM4001662	99. 250	58, 594	135, 514	19.947	25.792	43.075	21.822	28.747
	NT2RM4001666	108, 596	50.059	60. 195	8.757	26.897	80. 547	55. 639	44. 557
	HT2RM4001670 HT2RM4001682	23.010	37.857	52, 107	34:229	26.474	24.078	19.040	48. 902
	NT2RM4001710	71.974	22,009	43.652	12.553	17, 193	33.805	36.338	25. 346
	NY 2RM4001712	30.145	17. 963	29.768	6.775	12.959	13.705	17. 401	11.444
25	NT2RM4001714	39. 284	71, 253	45.168	23.590	23.852	34.014	32.992	44. 464
	NT2RM4001715	39.876	47.568	68. 485	29.814	28.676	29.317	23.694	38.125
	NT2RM4001727	18.826	16.671	24. 630	8.765	12. 534	14. 525	9. 624	7. 446
	NT2RM4001731	163.786	60.747	103.744	21.266	23.073	109.348	70.159	88. 870 48. 984
	NT2RM4001735	25. 147	42.977	27.836	33. 257	23.484	16.531	22.623 7.382	14. 863
	NT2RM4001739	29. 621	22.031	33.503	11.627	16.721	10.593 49.703	68, 739	91, 553
30	NT2RM4001741	117.616	80.979	99.834	34.861	34.797	37. 464	33. 824	23. 274
	NT2RM4001746	61.847	44. 910	113. 561	21.148	25. 420	34. 023	22. 194	15. 154
	NT2RM4001754	72.161	34.709	70.656	13.473	10.724	21.161	24. 761	19.803
	NT2RM4001757	38.117	23.659	28. 972	12.593	12.781	14. 153	7.027	6.943
	NY2RM4001758	24.391	23.518	60. 158	17.044	37. 261	58. 428	34. 390	27. 280
	NT2RM4001768	51.099	53.221	13, 303	2.644	9, 532	9, 892	6. 237	4.050
35	NT2RM4001775	15.024 24.497	11.154 20.843	16. 325	5.116	12.075	8.815	13. 233	6.515
	NT2RN4001776	44. 218	34.754	35. 521	11.654	27.683	28.899	24. 397	19.284
	NT2RM4001783	75. 698	74.949	146.739	24.426	38.218	21.996	28. 324	24. 241
	NT2RM4001810	25, 287	22. 294	22.627	8.986	12.014	13.754	22.602	19.691
	NT2RM4001813	108.290	15.721	11.311	3.071		7.061	9. 406	9.278
	NT2RM4001818	55, 110	32. 332	35.827	10.603		30.893	25. 538	20.147
40	NT2RM4001819	221.187	103.477	118,661	33.955		117. 958	105. 557	45, 891
	NT2RM4001823	31.566	19. 207	30. 580	9.100		18. 948	23.046	12.498 20.662
	NT2RM4001828	33.606	37.243	60.904	39.892		52.576	22.264	33. 367
•	NT2RM4001835		48. 485	36.681	12.402		32.404	46, 063	49.677
	NT2RM4001836	68.101	53.948	86.019			44. 492	38. 346	53.737
45	NT2RM4D01841	77, 551	75.005	64.963			60.179 18.888	13. 674	12.515
45	NT2RM4001842		31.217	153. 53B		1 057	30. 666	19.358	23.477
	NT2RN4001843		47.021	41, 491					17,632
	NT2RM4001856		17.427	22.905					27.815
	NT2RM4001858		13.809						30.143
	NT2RM4001861		55.955 33.911						32.578
50	NT2RM4001863		38. 767						30.584
	NT2RM400186								110.541
	NT2RM4001869								26.533
	NT2RM400187							135.056	71.907
	NT2RM400188								
	MT2RM400188						40.932	20.399	40. 632
55	M1 EVM-00 100	32.000							

Table 75

	NT2RM4001889	44.826	54, 188	57.058	17.324	30.679	30.391	31.401	27. 309
			21.032	38.644	10. 368	15.617	23.290	26.653	24.028
	NT2RN4001894	***	37.135	42.706	11. 443	18.977	24.084	62.995	21.376
5	NT2RM4001897		37. 833	50. 793	22.892	10.010	19.933	39.828	71. 231
	NY2RM4001899			131.041	19.900	22.521	28.037	22.888	34.298
	NT2RM4001905			167.103	32.535	29.282	32.842	21, 101	29.820
	NT2RN4001922		18.972	11.870	12.179	5.722	7.704	2.893	19.882
	NT2RM4001930	9.761	9. 323	20.059	5. 226	22.340	8,605	6.836	2.737
	NT2RM4001938	13.300	28.342	53. 112	22.045	19.769	35.835	24. 329	24.211
10	NT2RM4001940	44. 499		137. 250	99. 314	68.782	123.550	44.362	143. 236
	NT2RM4001942		67.064	218.754	37. 265	39, 359	37. 249	28.374	31.774
	NT2RM4001953	73.750	33.648	57.473	21.916	18, 921	11.704	7.776	32.933
	NT2RN4001965		24.684	41, 501	12. 421	18. 343	29, 179	21.379	18.604
	NT2RM4001966	49.431	22.954	33.007	12.456	14.747	23.958	15.690	13.553
	NT2RM4001969	82.202	23.827	35. 591	10.813	20.091	38. 983	35. 402	27. 290
15	NT2RM4001974	50.759	32.744	64. 327	26,669	29.268	32.957	29. 294	45. 426
	NT2RM4001979	64.506	28. 217	65.730	29.832	30.129	51.434	39.037	38. 269
	NT2RM4001980	8.940	10.121	18. 976	9. 204	7.020	7.587	10.490	17.931
	NT2RM4001984	76.782	27.2191	64.310	10.713	13.598	56.046	41.155	21. 341
	NT2RM4001987	19.064	9. 935	20. 167	9. 513	9. 423	13.449	15.551	64.982
	NT2RM4002018	23.330	15. 361	28.649	4. 482	9.866	15. 203	14.895	11. 409
20	NT2RM4002033	103.629	76.058	255.894	33.739	36.068	40.994	22.684	32.604
	MT2RM4002034	97, 025	74.014	204, 281	25. 591	40. 356	66.335	30, 838	29.885
	NT2RM4002044	128. 284	97.260	283.326	56.682	49.448	68.685	42.993	58.693
	NT2RM4002047	42.016	31.010	47.604	17. 496	19.793	15.043	24. 593	16.651
	NT2RM4002054	75. 334	24. 437	33, 919	5. 362	20.426	36.508	26.858	12.455
	NT2RM4002055	28.223	41.574	41, 231	17,667	21.073	24. 192	30.052	56.881
25	NT2RM4002059	24.790	47.792	30.688	32. 255	11.889	26.659	17.375	42.684
	NT2RM4002061	15. 353	22.159	24.342	33. 358	8. \$69	13.680	9.654	12.890
	NT2RM4002062	35.603	17.782	25.712	9. 437	13.693	23.679	11.468	12.877
	NT2RM4002063	106.902	59. 539	161.049	27.157	37. 323	44.770	45. 190	17.589
	NT2RM4002066	69.187	29. 278	44.089	14.142	12.777	29,901	23.625 19.168	38. 472
	MT2RM4002067	72.915	65. 950	164.446	33.322	23. 243	19.453	13, 427	15. 358
30	NT2RM4002073	26.509	19, 553	24. 129	7. 501	12. 225	9, 267	19.038	9. 923
	NT2RM4002074	23.768	16.727	27.356	9. 430	10. 288 8. 179	19.921	8.913	5.764
	NT2RM4002075	14.729	8, 566	14.082	6.113	12.370	22.729	21.957	5.088
	NT2RM4002076	33.772	34. 570	24. 768	12.754 29.244	28.319	38.890	38.136	27.441
	NT2RN4002078	65.837	45. 074	59.931	29.519	33.925	46.864	32.277	29.982
	NT2RM4002081	72.328	49. 374	162.917 24.293	4. 626	7.828	18.917	11.824	4.512
<i>35</i>	NT2RN4002082	31.523	20, 963 12, 906	28. 190	14.073	16. 132	8.993	10,746	15.942
	NT2RM4002093	13.703 48.477	33.601	44. 587	16.373	19.020	42.752	31.367	24.718
	NT2RM4002109 NT2RM4002115	52.087	16. 294	25, 726	5.046	11.691	15.294	19.312	5.666
	NT2RM4002118	6. 461	10. 205	16.364	2.841	6. 221	5.928	9. 423	8.612
	NT2RM4002128	24.014	12.586	38.670	8, 609	8.704	17.808	16.887	18.787
_	NT2RM4002137	60.650	30.735	54.930	9.746	20.827	30.629	27.756	30.682
40	NT2RM4002139	59.820	72.323	217.660	35.299	32.433	22.926	18. 198	31.925
	NT2RM4002140	61.939	27.988	54.095		18. 951	36.147	28.930	19.620
	NT2RM4002145	55.935	18.752	37.184		24. 220	25.455	54.028	17.830 22.927
	NT2RM4002146	10.714	7.232			4, 463	6.475	3.969 8,266	7.504
	NT2RN4002161	21.929	10.374			7.983	12.456	10.913	19.766
AE	NT2RM4002174	36.217	21.020			14, 155		18, 490	38, 988
45	NT2RM4002178	51.201	34. 975			26.852			41.037
	NT2RM4002180	88. 245						35. 104	
	NT2RN4002185	60.374	34.725		10, 870			_	
	NT2RM4002189	443.685	125.746						
	NT2RM4002194	110.410	60. 176						
50	NY2RW4002198	19, 112	25. 320						
50	NT2RM4002205	86. 368	52. 183						
	NT2RM4002213	87.023	29.632						
	NT2RM4002216	28. 034	36.860						
	NT2RM4002226	59. 214	25.842						
	NT2RM4002237	84.115	47.30						
55	NT2RM4002240	21. 140	20.81	18.200	11.22	4.61	-1-,,,,,,,,,,		

Table 76

	NT2RM4002251	39. 895	25.621	38.004	9.808	12. 483	27.050	27.880	15. 570
	NT2RM4002256	62.880	50. 437	132.459	16.059	20.051	22.911	18. 973	36.148
_	NT2RM4002252	40.381	19.221	18.726	4.067	10, 643	11.552	18. 506	11.180
5	NT2RM4002286	33. 927	16.242.	29.395	7. 271	10.706	15.907	16.746	45. 558
		31.555	29. 432	34. 470	12. 227	15. 207	18. 832	24. 174	41.738
	NT2RM4002276	24, 493	44.932	54. 554	19. 947	24.631	19.085	14. 211	28.361
	NT2RM4002278	73.045	68. 535	120, 767	28. 971	77.810	35.833	33. 197	34.350
	NT2RM4002281	95. 529	67. 191	148. 977	16. 383	32.822	42.647	36, 149	22.550
	NT2RM4002287	37. 325		32.626	7.879	22. 188	17.681	21. 208	18.691
10	NT2RM4002294		40. 622	14. 186	6. 186	12.213	8.996	13. 334	20. 467
	NT2RM4002298	15. 253	25.056	24. 351	8.779	13. 453	11.537	16.605	21.093
	NT2RM4002301	25. 506	22. 524	40. 307	8. 697	15. 098	30.071	33. 558	17.520
	NT2RM4002305	64. 514	27.130	108.848	13. 787	18.840	15, 998	23. 739	23.002
	NT2RM4002323	46. 276	37. 334		13.787	61.856	67.867	63. 381	16.555
	NT2RM4002334	84.665	44. 953	240.849	4. 286	11.781	13.743	14. 276	7.602
15	NT2RM4002339	40. 226	15.664	17.738		14. 835	5. 571	6.021	.15.852
	NT2RM4002344	15. 209	14. 735	15. 127	5. 186	49, 725	20.214	15, 169	93.476
	NT2RM4002345	29. 537	16.084	44.040	7. 161	10.828	17.765	20. 622	16.556
	NT2RM4002352	25. 146	26.320	39.068	10.070		13.358	9.862	21.089
•	NT2RM4002362	22.727	18. 9671	35. 121	7. 780	16.102	15.723	21.117	10.960
	NT2RM4002373	49, 413	25. 049	39, 501	16. 293	10.820	25. 509	12. 233	16.554
20	NT2RM4002374	45. 312	17. 702	80.866	14. 495	13.876 15.635	33.518	17. 499	20.037
	NT2RM4002376	44. 035	32.785	33.965	15.793 56.564	36, 130	62.968	25. 071	60. 431
	NT2RM4002383	143.921	114, 177	338.801 23.593	13.554	0.000	15.764	10. 120	21. 189
	NT2RM4002390	19. 946 33. 574		55. 577	19.871	29, 143	36, 917	34. 014	15.071
	NT2RM4002398		85.078 25.690	44, 155	15.629	15. 274	43.916	36.612	24.609
	NT2RM4002409	62. 430	27.569	49. 085	13.732	29. 300	20.609	24. 789	22. 958
25	NT2RM4002414	122. 797	24.210	57.361	13. 303	21.819	19. 128	27.861	33. 288
	NT2RM4002438	60. 880 50. 958	29.949	58.790	16.516	17.087	22.853	27. 261	86.320
	NT2RM4002440 NT2RM4002445	85. 102	43.893	64.557	15. 166	30.454	59, 828	43.072	34, 360
	NT2RM4002450	29.806	50.782	20.662	10. 226	5. 031	56.095	6. 391	48.088
	NT2RM4002452	38. 119	24.046	27. 781	13.792	11.741	21.974	28. 908	14.192
	NT 2RM4002457	56. 998	45. 958	72.065	21. 106	21, 980	25.587	22.709	26.372
30	NT2RM4002458	17. 499	9. 159	12.416	3.859	12.704	4.423	1.834	7.476
	NT2RM4002460	37. 183	7, 502	15. 263	2.616	9. 265	20.827	12.805	1.464
	NT2RM4002464	12.680	10.529	5. 512	5.737	10.707	1.669	5. 391	12. 187
	NT2RM4002479	85.068	45.694	66.175	35. 340	44.661	52. 236	42.316	33.845
	NT2RM4002482	714. 577	349.138	482.476	135. 984	180.855	462.386	321.086	260.860
	NT2RM4002489	41.987	36.475	28. 303	18.347	20. 193	. 45. 527	22.970	15.427
35	NT2RM4002493	101.547	19.009	34.214	7.129	20.517	58.926	20.613	6.136
	NT2RM4002499	104. 508	114. 364	295. 841	132.961	45.496	125. 546	54.809	138.353
	NT2RM4002504	130. 575	85. 186	319.621	58.095	51.615	65. 385	43. 397	39.625
	NT2RM4002506	17. 534	7.716	22.097	8.307	8.641	11.973	11.217	19.715
	NT2RM4002510	20. 570	20.274	28. 261	7. 195	10, 108	9.354	16.982	8.405
	NT2RM4002527	29.097	14.199	26.008	7.215	11.820	15.320	15.507	11.537
40	NT2RM4002532	119.266	103.485	252.069	38.479	49.581	51. 534	30.506	48.759
	NT2RM4002534	46.720	29. 222	28.381	12.470	17.005	30.785	27.381	25.218
	NT2RM4002535	150.736	124.425	370.470	71.472	69.884	70, 122	44. 328	39.348
	NT2RM4002554	46.680	4.678	15.042	2. 434	7.853	8. 287	11.868	8.546
	NT2RM4002558	64. 523	30.756	60, 861	17. 849	28. 435	32.697	50.330	26.839
	NT2RM4002565	26.150	21.759	29.418	10.020	13.855	14.504	15. 952	20.143
45	NT2RM4002567	13.750	9, 555	16, 128	7.961	6. 533	14.816	11. 242	24.778
	NT2RM4002571	64. 981	32.370	51.874	13.381	25. 113	37.880	40. 593	30. 327
	NY2RM4002572	21. 932	17.415	44. 482		9.094	15.081	8. 955	11.463
	NT2RM4002577	13. 390	34. 537	17.827	8. 379	17.150	9.208	20. 440	135. 375
	NT2RM4002583	43. 872	21.818	41.335	7. 938	12.820	25. 087	16.879	8.165
	NT2RM4002584	48. 978	41.874	47. 589		15. 387	18.002	26.572	29.591
50	NT2RM4002593	43.140	21.408	34.068		17.846	27. 459	22.581	18.025
	NT2RM4002594	53, 494	32. 355	54. 474		23.934	38. 188	30. 209	28.918
	NT2RM4002604	49.799	31.218	31.584		10.658	52, 255	31.422	27.262 10.800
	NT2RM4002614	18.848	9, 948	15.663		10.103	19. 152 25. 551	15. 480 20. 412	22.945
	NT2RM4002616	52.378	28. 130		6. 189	16.589	28. 155	15. 957	8. 295
	NT2RM4002623	31.915	15.505			6.856	27, 565	17.040	9. 308
55	NT2RM4002634	27.202	13.607	23.468	4. 566	0.030	1 41.303	11.040	1 3.300

Table 77

				A 217 T	2 074	1 466 1	2.585	2.436	4. 543
	NT2RM4002636	2.342	5. 234	9.517	3.874	1.465			
	NT2RP1000002	114, 491	47.508	61.586	25.000	29.448	84.026	73.878	59. 624
	NT2RP1000006	71.057	28.511	44.224	10. 202	17. 523	40.868	37.373	15. 237
5			9, 953	16.089	4,506	3.649	7.738	3,651	8.661
	NT2RP1000015	7. 192		0.000	0.000	0.000	2.690	4.737	0.000
	NT 2RP 1000018	5.882	0.000					21.883	51,696
	NT2RP1000034	273.802	61.801	59.676	50.413	101.761	283.598		
	NT2RP1000035	14. 407	14. 328	5.278	5.331	3.905	19.347	5. 560	9.946
	NT2RP1000040	2.229	2,143	2.569	1. 482	0.842	0.251	1.226	0.963
		2.962	1.516	2.106	0.450	3,003	1.458	1.788	0.000
10	NT2RP1000042				1, 520	1, 452	2.742	0.779	17.389
	NT2RP1000048	3. 312	5, 643	4.404		7.544	14. 598	18. 930	19.749
	NT2RP1000050	37.260	7. 381	21.735	7.969			1.919	18.856
	NT2RP1000056	2.575	8.244	12.209	2.506	2. 248	50.055		
	NT2RP1000058	7,701	2.152	6.853	1.889	5.740	5.703	5.884	4.654
	MT2RP1000063	17.863	6,661	8.488	2.745	0.000	7. 494	5. 484	2.401
			5. 197	4, 140	0.833	1.697	1.068	0.863	1.468
15	NT2RP1000068	4.612		72.321	37.376	27, 104	99.463	69.787	134. 954
	NT2RP1000072	143.838	99.413				0. 523	4. 055	9.765
	NT2RP1000073	1, 552	1.742	0.000	0.919	0.996			3. 421
	NT2RP1000078	2.896	0.000	0.000	0. 230	0.741	0.763	0.567	
	NT2RP1000079	49.027	29.65%	15.514	6.677	6.650	9.256	18. 182	28.375
	NT2RP1000080	16.385	13.693	8.875	4. 934	5.832	9.673	15. 737	12.194
			3.761	10.248	2.946	7.423	5. 286	3.826	0.000
20	NT2RP1000086	7. 169				3.506	2.887	0.000	2.053
20	NT 2RP 1000087	0:000	5.038	0.000	1. 221			0.418	13.867 -
	NT2RP1000089	4.302	9.012	-8.097	5.674	2. 992	4.624		
	NT 28P1000090	52, 428	58.867	69.998	38, 821	17. 374	29.637	35.043	79. 235
	NT2RP1000100	3, 207	3,774	1.540	2.138	1.112	1.149	0.000	1.791
	NT2RP1000101	92,707	46.496	68, 186	33.782	33.861	36.104	55. 994	56.718
		4. 451	9. 940	6.651	2,623	8.151	2.765	11,052	2.965
25	NT2RP1000111			0.000	2.480	0.000	1.727	2,041	2.374
23	NT2RP1000112	3.985	3.478			2.,553	12.703	2.802	42.544
	NT2RP1000124	24. 505	9.928	6.917	5.644		62.060	44. 484	52.427
	NY 2RP 1000125	24.817	79.995	139.555	49.819	97.770			
	NT2RP1000129	28, 170	30.324	26.037	10.799	3.638	16.350	16.315	13.950
	NT2RP1000130	5. 381	7.279	14.556	2.578	10.778	12.987	0.000	20.710
	NT2RP1000154	17.054	18.625	18.032	7.765	17.883	13.855	12.502	19, 133
20			7.739	9.822	4.142	3, 589	2.512	6. 952	17.030
30	NY2RP1000163	18. 531			1.557	5. 549	3.844	7.224	15,609
	NY2RP1000170	14.775	6.603	3.911			1. 497	5.060	0.857
	NT2RP1000174	10.066	4.006	4.875	1.601	3.951			56.201
	NT2RP1000181	108.209	58.429	137.843	40.129	31.719	74.897	73.935	
	NT2RPTOCO191	9, 285	6.645	5.460	3.099	6.842	12.524	5.864	2.766
	NT2RP1000202	4.547	3.462	7.203	6.298	6.151	3.022	2. 481	4. 122
0.5		0.000	0.000	4.313	1.852	1.396	1.558	2. 101	1.136
35	NT2RP1000239		5.330	3.864	1, 538	6.834	4.100	5. 184	5. 579
	NT2RP1000243	10. 228		2.512	1.848		2.012	5.711	5.678
	NT 2RP1000255	6.844	3.187				8.515	4.509	4. 367
	NY2RP1000259	10.073	6.510	10.276	1.573			1.763	0.000
	NT2RP1000261	0.000	0.000	0.000	0.000		0.000		
	NT2RP1000269	233, 453	119.331	130.392	48. 933		111.105	129. 953	95. 341
40	NT2RP1000271	504. 212	314.887	684.003	191.587		351.080	721.963	268. 189
40	NT2RP1000272	130, 317	52.877	78.345	38.313	30.575	71.135	50.465	37.296
	NT2RP1000279	103.540	36.699	55. 522	23.329		68.415	50.629	9.388
				295. 250	136. 106		257.258	215. 344	195.667
	MT2RP1000290	383.695	214. 173		43.735		85.003	75. 569	61.144
	NT2RP1000293	139, 263	71.666	91.679				105. 443	25.701
	NT2RP1000300	219. 317	94, 497	120.961	62.228		166.238		54.085
45	NT2RP1000324	205. 212	96.463	109.241	73.482		120.952	75.697	
45	MT2RP1000325	567, 975	208.141	235. 225	74.690	106.786	298.190	175.163	181.979
		114. 548	37.978	60.587	21.766	22,713	70,707	48.865	22. 186
	NT2RP1000326			12.198	9.94		9.595	5, 409	16. 164
	MT2RP1000331	14. 215	11.082					80, 360	48.737
	NT2RP1000333		62.474	124.398	35.732			5. 855	5. 234
	NT2RP1000336	5.071	3.476	0.000					
co.	NT2RP1000347	8.732	4.239	0.000				4. 829	4. 180
50	NT2RP1000348		3.224	2.495	2.89			4.511	1, 450
	NT2RP1000349		4. 441	0.000			3.407	3.025	2.512
			80.510					33.300	118.905
	NT2RP1000353								110, 239
	NT2RP1000356		45. 385						87.310
	NT2RP1000357	213. 820	128, 901	421.667					67.426
E E	NT2RP1000358		64.055	108. 939	32.77	8 41.72	110.904	74.510	1 91.420
55									

Table 78

									00 704
	NT2RP1000360	297.314	134.601	191.999	71.819	85. 890	202.062	147.810	89.594
	NT2RP1000363	364, 040	212.933	280.442	136.437	123.748	247.266	256.906	128.344
	NT2RP1000376	127.768	49.154	84.631	29. 920	40, 910	71.095	82.258	43.951
5				56.520	52. 245	252, 336	185. 039	121.336	65. 534
3	NT2RP1000386	39. 353	145.725					2.424	3.461
	NT2RP1000407	2.663	0.197	0.000	2. 423	0.000	3.032		
	NT2RP1000409	0.000	5.878	0.000	0.850	0,000	0. 424	0.000	0,000
	NT2RP1000413	7, 153	2.048	2.681	0.000	8. 303	4.015	0.344	0.307
	NT2RP1000416	0.000	0.000	0.034	0.000	0,000	0.000	0.000	0.000
				8,733	3, 988	5, 668	7,649	7, 116	4, 283
	NT2RP1000418	9. 174	4. 984				0.000	0.147	0.000
10	NT2RP1000420	2. 125	0.924	0.000	0.000	0.000			
	NT2RP1000434	0.000	19.791	0.000	0.750	0.000	0. 189	1.654	0.000
	NT2RP1000439	134. 853	56.272	115.668	51.887	49. 782	73. 229	64.079	15.355
	NT2RP1000443	58. 432	1.440	0.000	3. 540	5, 276	7, 299	4. 900	2.656
			2.955	0.800	3, 240	1.187	3, 303	1.052	3.063
	NT2RP1000447	3. 820				1.043	0.314	0.856	0.000
	NT 2RP1000448	3.888	0.697	0.000	0.778				3, 138
15	NT2RP1000451	5. 766	4.110	3.245	4. 480	1. 272	3. 036	1.022	
	NT2RP1000458	277, 437	139. 151	249.632	114.073	87, 709	243.919	188. 141	160.796
	NT2RP1000460	216, 381	129.722	192.470	86.161	96, 273	135. 913	170. 172	91.267
		290.518	221.955	402.881	192.151	210.010	230. 322	182. 401	205.887
	NT2RP1000465			61.862	19, 161	13.854	16.791	11.220	11,713
	NT2RP1000468	29. 203	30.933					101.037	71.927
	NT2RP1000470	247. 991	94.630	118.548	33.073	62, 185	113. 536		
20	NT2RP1000477	3. 039	1.894	0.000	0.887	1.636	2.721	1.261	1.757
	NT2RP1000478	2.842	0.655	0.000	0.363	1. 122	0.412	1.375	0.000
	NT2RP1000481	5. 676	0.693	1,376	2.294	1. 991	0.993	2.480	1.941
		5, 004	0.820	0.000	1.070	0.687	1. 252	0.401	0.344
	NT2RP1000493		62.178	133.983	29. 869	42. 569	122. 982	62, 701	55. 329
	NT2RP1000513	183. 214					110. 978	93.419	\$2.294
	NT2RP1000522	183. 947	57.483	120.005	32.529	32. 275			
25	NT2RP1000533	21.686	8.198	15.700	5,816	6.071	12. 902	9.030	5, 528
20	NT2RP1000544	3.732	10.988	1.704	2.465	2. 581	6.543	9. 371	6.069
	NT2RP1000547	0, 300	0.310	0.170	0.000	0.000	0.000	0.000	0.000
	NT2RP1000551	3.715	1.322	3.371	0.657	1.870	1.149	3. 287	1, 199
				7, 630	1.128	0.978	9.115	8.337	2, 192
	NT2RP1000567	18, 148	4. 535				2.846	0.662	0.000
	NT2RP1000574	2.807	2.740	4. 159	0.000	1.266			
30	NT2RP1000577	5.767	6.059	6.234	2.033	4.066	4.517	1.545	3, 168
30	NT2RP1000579	13, 591	6.812	7.808	2.066	3. 452	4.699	7.020	6.279
	NT2RP1000581	23. 446	8.664	15. 950	5. 531	6.046	15. 075	12.761	9.085
	NT2RP1000593	6.058	14. 376	5.780	2.580	5. 057	9.162	5. 483	15.975
				5, 413	5. 134	3, 748	4.785	3.835	2.255
	NT2RP1000604	3.081	4. 126				13. 382	13.762	3,825
	NT2RP1000609	27. 487	3. 174	10.612	2. 728	3, 986			
	NT2RP1000613	4. 356	2.265	1. 529	1.001	0.000	1.184	2.710	0.767
35	NT2RP1000622	15, 005	7.496	8.013	1.968	1, 752	7. 985	7.518	6.485
	NT2RP1000627	17, 344	14.772	22,410	6.441	12.047	16. 356	20.729	10.336
	NT2RP1000629	15.718	4.144	12.352	4.104	4. 312	7.820	11.024	7.693
			32.499	52,699	15, 138	14, 415	30.508	33,741	18.936
	NT2RP1000630	65. 249				10.606	19.791	14.683	16,200
	NT2RP1000639	43. 900	18. 204	18.020	10. 187			17. 026	76.883
	NT2RP1000640	86. 217	156.971	37.078	60.057	32.726	29. 102		
40	NT2RP1000646	7.394	16.894	13.629	5. 542	5.660	7.382	1,582	2.851
	NT2RP1000659	26.494	13.979	53. 935	11.276	9, 11,9	12.945	10.602	15.936
	NY2RP1000674	10, 820	5. 502	9, 633	4.224	4. 542	3.907	5. 942	5.755
	NT2RP1000677	187. 310	76.173	99.589	25. 959	49.679	90.146	95. 230	63.227
			5. 907	7.263	2.229	1. 965	2,520	3.853	6.223
	NT2RP1000679	9. 839					17.736	20.738	18.729
	NT2RP1000688	30.741	21.137	41.993	9. 852	14. 205			
45	NT2RP1000689	8. 594	2.814	13.021	1. 222	4. 171	7. 394	4, 473	3. 167
	NT2RP1000695	1.813	3. 104	2.068	0.810		0.000	0.786	0.000
	NT2RP1000701	1.280		0.000	0.000	0.000	0, 855	0.000	0.607
	NT2RP1000702	4. 112		8. 473		1.698	1,616	4.749	0.000
						0.000			0.000
	NT2RP1000713	0. 233							76.919
	NT2RP1000721	199. 987				64. 142	102.872		
50	NT2RP1000730	24. 414	16.302	64.370		6. 129		8.948	
	NT2RP1000733	9.992	13.894	13.138	3.593	3.087	6.945		
	NT2RP1000738	357. 551					211.940	169.539	140.421
	NT2RP1000739	261. 372							
	NT2RP1000740	60.717							
	NT2RP1000746	13. 275	9, 551	20.132	3.376	1.635	3, 601	3.265	1 3.309

488

Table 79

	NT2RP1000750	134.663	52.958	80.346	28.605	36. 158	71.713	92.250	39.685
	NT2RP1000751	17,717	44. 325	31.941	32.295	15. 461	19.059	18.084	64.708
_	NT2RP1000767	12.860	6. 572	9.057	2.510	3.872	3. 120	5.111	4.085
5	NT2RP1000769	27.412	21.636	18.089	7.324	7.758	13. 441	12.436	7.317
	NT2RP1000780	7.664	2.995	3.269	2.715	2.030	0.000	0.000	0.000
	NT2RP1000782	11.618	23. 259	28.607	5.885	16.596	14. 946	5. 301	7.061
	NT2RP1000796	118. 585	56. 532	75.809	15.096	41.498	78.341	73.407	26.885
	NT2RP1000797	215.680	107.927	100.844	28.806	53.841	131.952	306.946	77.792
10	NT2RP1000800	5. 249	3. 787	2.211	1.617	7.056	3. 306	3. 512 24. 570	2.851
,,	NT2RP1000825	49.312	22.623	29.009	4.529	15. 271	16. 815 25. 255	29. 305	12.101 27.307
	NT2RP1000833	67.848	23. 702	41.132	9.260	13.328	19, 117	14. 348	17.998
	NT2RP1000834	21. 157	17. 555	15.686	11.112	2.621	7. 219	5. 827	5. 382
	NT2RP1000836	12.434	11.272	7.839	3.196	23.029	41. 395	35.068	27.483
	NT2RP1000837	98. 743	40. 415 11. 209	35. 656	21.833 4.957	5. 131	7, 919	3.229	5. 512
15	NT2RP1000846	14.775 27.431	18. 237	16.588	10.757	10, 320	14, 784	19. 182	10.029
	NT2RP1000847	214.374	87.847	128.937	45, 113	51. 955	144, 598	108.723	51.968
	M12RP1000856	26. 023	29.514	67.757	23.663	28, 185	38. 015	15.874	11.458
	NT2RP1000860	163.711	61.1001	101.078	35.949	41.953	87.889	80.204	48.859
	NT2RP1000902	24.271	31.899	49.716	12.862	11.237	22. 189	17. 326	17.501
	NT2RP1000903	68.716	24. 490	31.806	9.135	15. 239	68. 242	28. 337	14.115
20	NT2RP1000905	25. 662	13. 385	22.530	7.568	3_894	6. 452	12.011	13.929
	NT2RP1000915	22.768	32.699	39.412	17.920	10.752	16.493	10.059	19.431
	NT2RP1000916	36, 356	17.076	24.787	8.241	2.752	29.963	18. 336	4. 134
	NT2RP1000921	20.200	10.536	22. 363	8.324	5.717	15. 896	15. 473	4.407
	NT2RP1000943	9. 440	4.278	14. 836	10.665	2. 682	3. 445	1.686 29.682	1.791 34.740
	NT2RP1000944	65.067	27.816	40.730	12.441	18. 584	43. 858 17. 723	9. 687	20.200
25	NT2RP1000947	18.414	12.386	22.697	15. 197 5. 358	10.849	20. 625	13. 192	15.554
	NT2RP1000954	28. 307	24. 912 38. 788	28. 425	23.030	11. 285	20. 525	21. 953	28.920
	NT2RP1000958	21.987 84.562	81.956	164. 902	59.895	32.501	60, 329	47. 308	77.704
	NT2RP1000959 NT2RP1000966	104. 461	73.705	101.907	58.853	28.479	65. 560	39. 891	37.125
	NT2RP1000974	213.892	124, 166	171.079	71.813	73.877	160.514	104, 131	41.698
30	NT2RP1000980	16. 802	11.080	6.958	4.146	7.799	7.626	6.311	2.017
50	NT2RP1000981	50, 385	24. 506	35.067	13.841	17.653	24, 416	15. 302	5.946_
	NT2RP1000988	19.623	11.058	22.054	9.003	7.658	18.310	15. 545	11.394
	NT2RP1001002	56.891	33.510	22. 993	6.717	20.078	27. 348	21.988	16.177
	NT2RP1001004	23.268	13, 134	13. 405	6.295	5.883	11.999	12.399	18.783
	NT2RP1001007	29. 127	10.102	12. 426	8.003	3. 193	18.313 28.628	13.582	9.737 12.589
35	NT2RP1001011	36.507	27.547	42.002	16.657	13.048	29.607	9.620	52.526
	NT2RP1001013	9. 942	14.082	54. 179 30. 913	10.101	11, 200	9.468	17.655	12.776
	NT2RP1001014 NT2RP1001020	39.078	9. 107	36. 274	3.816	13.500	15. 563	15. 121	4.580
	NT2RP1001023	5309.613	985.566	1698.618	284.967	1874. 160	4332.654	3092.785	808.260
	NT2RP1001027	73.098	53. 184	34.629	18.681	24. 296	93.325	67.199	51.245
	NT2RP1001031	6.727	3.944	1.413	2.625	2.583	4.462	2.652	2.043
40	NT2RP1001033	34. 383	18.547	52.827	11.061	12.794	15.798	10.825	16.802
	NT2RP1001042	16.664	10.042	32.855	18.106	26. 513	10.262	8. 945	11.915
	NT2RP1001045		33, 846	51.766	24.186	48.474	72.682	35. 437	30.767
	NT2RP1001073		10.612	7.850	6.640	5.048	9.855	6.935	5. 520 16. 408
	NT2RP1001079		71.311	176.776	25.199	28.090	18.812	18. 139	11.376
45	NT2RP1001080		23. 422	19.061	11.316	14, 731	9.872	5.861	3.904
43	NT2RP1001113 NT2RP1001159		5.617 59.111	8. 219 125. 441	72.993	66.677	187.780	55,003	98.072
	NT2RP1001173		13. 137			17.090	13.269	9. 476	11.252
	NT2RP1001176		10. 035			14.457	10.458	9. 085	4.024
	NT2RP1001177		25.797			13, 900	29.445	22.230	7.579
	NT2RP1001185		76.839			27.738	39.654	27.055	27.069
50	NT2RP1001199		17.518			14.093	14. 390		11.780
	NT2RP1001209		19.355		18.438	19.648	28. 439		36.255
	NT2RP1001215					11.039	25. 483		15.808
	NT2RP1001225		20.250				26. 429		22.194
	NT2RP100124		8. 531				12.906		9.166
	NT2RP100124	6. 228	6.100	7.648	1.747	1.022	2.368	3.698	2.028
55									

## Table 80

								AVE T	47 637
	NY2RP1001248	49. 226	25. 943	116.548	10.461	11.820	12.652	13.256	17.837
	NT2RP1001253	16.172	14.468	19.494	5.712	7.057	20.880	11.966	15.830
_	NT2RP1001286	31.909	17.523	37, 293	9.003	10.973	24. 180	18. 180	18.610
5	NT2RP1001294	25.024	26. 137.	.24.014	7.577	12.732	16. 248	11.737	14.676
		20.570	17.865	14,990	7.914	7.089	11,711	10.424	6.370
	NT2RP1001302			61.003	20. 191	35, 975	42. 746	31,795	30.891
	NT2RP1001310	73.669	50. 596					43.729	26. 945
	NT2RP1001311	107.757	35.881	46. 474	17.712	21.645	48. 944		
	NT2RP1001313	55. 324	32.674	63.966	13. 492	14.367	18. 129	17.116	14.648
10	NT2RP1001324	35. 171	18.577	22.653	7.819	11.963	16.113	15.675	21.371
10	NT2RP1001349	44. 453	17, 959	25. 475	6.766	11.881	22.818	27.028	20.116
	NT2RP1001361	55.753	27.902	58.131	21.682	28.045	60.728	52.605	27. 148
	NT2RP1001379	126.769	137, 614	71.862	24.018	47.600	154.003	231.914	35.839
			89. 642	123.622	19.403	22. 929	45.989	34. 307	19.045
	NT2RP1001385	74. 494		24. 575	7.512	17.756	24. 165	18.832	16. 437
	NT2RP1001395	45. 302	31.340		12.632	9. 318	21.843	13. 537	8.295
15	NT2RP1001410	23.514	23.629	40. 104		4, 204	7. 482	8.833	-25. 347
	NT2RP1001424	10.618	33.112	10.799	2.636				
	NT2RP1001432	12.466	40. 995	9. 503	1.789	6. 323	5. 098	8. 187	7. 252
	NT2RP1001449	55. 536	20.728	66.767	10.440	26. 188	27. 184	29.004	30.274
	NT2RP1001457	30. 322	32.7213	37.777	8.330	12.956	20. 340	25.841	17.849
	NT2RP1001459	88.712	62.417	75.498	27.541	35.602	62.144	51.183	51.852
	NT2RP1001466	16.844	23.355	27.785	10.621	12.274	14.384	8.050	13.792
20	NT2RP1001475	89.839	111.813	276.258	35.857	23.078	34.083	16.906	15.713
	NT2RP1001482	9. 804	7. 238	3. 123	7.419	2.367	3, 451	2.538	1.692
				15, 730	1,433	3.642	8, 911	7,609	6.956
	NT2RP1001494	18. 452	17.405		2.456	0.000	0.086	0.162	0.765
	NT2RP1001500	2. 143	2.318	3.634					6. 297
	NT2RP1001517	14.740	13.801	16.801	3.704	5. 628	8. 123	9.615	
	NT2RP1001540	50. 226	35.070	52.423	11.150	17.869	36.090	28. 195	7.025
25	NT2RP1001543	87.779	27.665	55.068	12.390	25. 264	48. 623	28.462	18. 547
	NT2RP1001546	51.476	99. 185	143.880	25. 320	72.799	104. 259	38.212	42.007
	NY2RP1001550	67.741	63.428	53.684	15.107	31.309	40.950	26.433	16. 133
	MT2RP1001553	34. 956	17, 566	22, 966	10.039	10.915	17. 357	20.710	19. 945
	NT2RP1001555	33. 240	52.576	54,908	25.406	21,523	42.121	29.401	24.807
	MT2RP1001563	30. 536	23. 522	26.745	10.623	16.136	20. 228	17.699	11.340
30	NT2RP1001569	90.271	31.802	37.662	7.791	18,755	32.159	31.572	22.545
30					20.979	38. 153	69.983	85.177	68.021
	NT2RP1001584	125. 503	64.642	101.860		9. 376	19.848	14.150	13.608
	NT2RP1001599	25. 536	22.635	29.822	7.141			14.067	12. 210
	NT2RP1001616	38.077	18. 321	20.981	7.268	5. 256	12.873		
	NT2RP1001654	77.215	24. 275	26.850	14.308	14.684	36.754	26. 803	17.786
	NT2RP1001665	20.132	15. 451	16.433	5.156	9.958	5. 979	8.761	8. 109
35	NT2RP1001679	261.384	264,730	245. 821	192.156	85.798	197.731	172.668	434.739
00	NT2RP1001681	21.960	21.892	16.974	17.231	5. 379	21.608	10.982	20.811
	NY2RP1001694	27.832	32.368	36, 517	12.438	29, 150	109, 147	231.086	69.267
	NT2RP2000001	79.348	34.825	26.858	8.546	17. 504	24.165	27.629	18.039
	NT2RP2000006	32.218	26, 701	47.407	11.066	8.723	14.994	13.215	12.652
		54. 262	32.503	34, 116	12.829	11.972	20.410	21.705	11.281
	NT2RP2000007			59. 562	20.809	17. 226	20.509	17. 286	54, 391
40	NT2RP2000008	34.810	31.036			5. 341	8. 149	10.076	5.865
	NT2RP2000010	12.320	9. 820	24. 557	3.019			50.108	45.745
	NT2RP2000011	121.718	115. 419	216.553	41.153	44.035	64. 567		
	NT2RP2000027	74.085	69.757	136.369	23.981	28.217	40.308	24. 108	20.710
	NT2RP2000028	23.699	28. 386	27.077	10.607	11.433	22. 532	14. 265	11.554
	NT2RP2000032	10.199	6.568	16.529	6.282	6.462	9.523	8.119	8. 527
	NT2RP2000040	383.423	222.501	199.099	79.455	81.787	229, 220	181.239	162, 128
45	HT2RP2000042	97.011	62.254	67.577	29.525	13.003	45. 921	45.196	41.158
	NT2RP2000045	73, 700	49. 722	66.899		17, 180	32.492	32.785	35, 403
		37, 323	46. 342			13. 292	43.534	29.174	17.962
	NT2RP2000051			69. 945		22. 707	40.001	40.807	38,782
	NT2RP2000054	99.806	54.072				26.794	25. 564	25, 156
	NY2RP2000056	57.518	40. 207	41,868		24. 303			
	NT2RP2000057	156,050	177.739	178.741		76.886	130.744	163. 333	207.593
50	NT2RP2000067	59, 366	13.414	39.371			22.699	22.699	5.023
	NT2RP2000070	107.618	50.674				83.478	48.688	26.235
	NT2RP2000076	48. 409	27.260	29.570			32.852	11.701	13.485
	NT2RP2000077	94.993	53. 327			14.024	49, 100	33.647	31,168
	NT2RP2000079	62.685				26.739	30. 432	16.329	18.678
	NT2RP2000088	71.164					42.871	35.756	8.836
55	IN THE EDGOOD	1	1 23.34.						
JJ									

Table 81

			** *** T	25 266 1	17 150 1	14. 253	18.714	15.927	10. 202
	NY2RP2000091	39.115	38. 293	35. 366	17.159		55. 591	34.478	54, 330
	NT2RP2000092	75.001	89. 256	171.691	60.810	53. 472		17.851	11.653
5	NT2RP2000097	31.201	13. 401	27.451	11. 261	15.139	18. 293	7.052	5. 446
9	NT2RP2000098	26.707	11.006	13.971	6. 330	7.991	11. 945		92. 458
	NT2RP2000108	169.612	134, 647	385.078	90. 234	79.343	81.573	54. 191	
	NT2RP2000114	32.814	21, 256	23.561	8.385	6. 127	16.427	11.227	18.744
	NT2RP2000116	24. 247	26.308	35. 305	21.085	8. 128	21.812	11.292	29. 326
	MT2RP2000119	87,773	75.708	213.188	30. 879	26.975	32.244	18.663	23. 323
	MT2RP2000120	28. 158	40.341	40.702	11. 423	17. 144	20. 974	18.758	14. 232
10	HT2RP2000126	68, 253	51. 174	75.714	25.719	32.146	30.674	19.806	13.086
	NT2RP2000133	40.974	21.405	31.855	9.468	16.094	19: 158	19.716	9.703
	MT2RP2000147	121.104	61.190	75.784	23.438	33.839	75. 147	46.430	37.718
	NT2RP2000153	96. 598	63.476	66.144	23.377	31.821	72.069	43.415	32.773
		115, 309	87.737	200.582	37.008	35, 422	38. 443	28.450	20. 252
	NT2RP2000156	24. 318	18, 095	28.697	14, 121	12. 284	22.086	12.179	10.763
15	NT2RP2000157		12.679	24. 575	5.678	7. 191	9.079	8. 105	9.807
	NT2RP2000161	9, 493	14. 646	19.908	3.979	5. 383	6,466	8.554	3. 206
	NT2RP2000168	11.413		150.036	37.188	58.850	114.315	90.491	66.465
	NT2RP2000173	228.420	98.033	71.096	15.404	30.614	50. 131	40.431	40. 206
	NT2RP2000175	78.839	44.514		14. 454	19.558	28.068	22.439	16. 249
	NT2RP2000178	60.513	42.174	41.614		44.541	64. 271	60.391	53.828
	NT2RP2000183	120.139	90.798	139.074	34. 168		45. 934	28.749	18.697
20	MT2RP2000195	91.304	70.037	204.874	30.805	27. 133	154. 895	68.553	248. 768
	NT2RP2000204	91,419	106.652	263.856	91.981	356.822		7. 227	6.812
	NT2RP2000205	30, 577	27.665	61.321	18.312	9. 596	17.099	31.136	31. 123
	MT2RP2000208	53. 204	48.346	85. 459	22.464	20. 371	37.407		
	NT2RP2000224	69.062	δ2.644	64.951	28.002	14. 265	42.146	33.510	51.634
	NT2RP2000230	56. 320	38. 161	51.891	19.712	16.865	28. 186	30.382	25. 164
25	NT2RP2000231	237.426	115.377	160. 416	68.560	85.769	155.055	119.086	87. 184
20	NT2RP2000232	49.708	32.849	24.700	10.366	11.881	31, 935	21.623	13.775
	NT2RP2000233	74, 158	43.941	52.603	20.024	20.149	47.211	52.894	52. 273
	NT2RP2000239	32,380	15.399	30.197	8.574	4.025	17.013	20.268	23.735
	NT2RP2000240	49.173	38. 363	78.202	15.737	15.654	21.302	17.214	20.486
	NY2RP2000248	17, 308	13, 339	13.368	4.823	12.687	8.493	12.992	9. 218
	NT2RP2000256	37.650	25. 977	25.477	12.706	9. 212	23.055	14.601	18. 126
30	NT2RP2000257	69.335	66, 181	244.979	45, 881	37. 192	46.969	31.322	46.624
	NT2RP2000258	39, 114	41,740	49.525	15.968	19.509	29. 341	17. 580	17.049
	NT2RP2000261	46.051	30.214	48.737	10.438	13. 441	22.674	19.894	19.556
	NT2RP2000270	73, 075	55.962	155, 102	33.557	25.014	49. 469	26.505	41.022
	NY2RP2000274	15.514	7.310	20.284	4, 327	6. 428	13.479	7.807	4.833
	NT2RP2000277	12.320	12, 198	8.692	2. 395	5.097	7.436	9.834	3. 452
35	NT2RP2000279	12.294	6.735	9.825	2.486	5.467	4.265	7. 545	6.898
	NT2RP2000283	63.324	49, 998	59,636	18, 166	19. 261	33.586	39.787	48. 270
	WT2RP2000288	38. 289	22.877	35.809	11.594	14. 150	24.632	25.978	24.657
		51. 997	39, 352	53.601	14.746	19.914	36. 153	31.476	28.603
	NT2RP2000289		71.227	206. 854	45.839	34.290	40.991	22.703	77.905
	NT2RP2000297	76. 236 28. 739	29.954	34. 444		10. 562	21.620	14.607	21.804
40	NT2RP2000298	29.075	14.696	16, 125		10.245	16.627	19. 121	11.456
40	NT2RP2000310		16, 201	24.879		13.651	24.922	17.858	30.618
	NT2RP2000327	45. 414 36. 600	35. 521	50, 933		23.798	33.981	22. 925	32.863
	NT2RP2000328			14, 112		11.584	34.848	35. 626	29.436
	NT2RP2000329	45. 820	29.353	89. 382		13.127	27.377	10, 155	15.517
	NT2RP2000333	33.894	26. 367			6. 291	12.498	8.041	5, 971
	NT2RP2000337	14.768	17.723			13. 030	26. 358	27.011	31.395
45	NT2RP2000346	53.051	82. 391	46.420		7.733	16. 593	11.667	6. 892
	NT2RP2000357	30.149	22.042	7 4 707			11.789	8, 150	7.184
	NT2RP2000358	16. 228	10.853	14.700				37.686	6. 290
	NT2RP2000366	82.288							12. 275
	NY2RP2000369	21.429					9.148	7.691	
	MT2RP2000376	205.303						139.412	79.987
E0	NT2RP2000394	31.766	23.882					24. 285	20. 279
50	MT2RP2000396	231.332	142.481					153. 536	
	NT2RP2000412							32.657	39.766
	NY2RP2000414		86.021						
	NY2RP2000420		33, 139					19.531	
	NT2RP2000422			24.54	8.16	6.449	11.697	14.485	17.945

491

Table 82

	Umannada dae	114 626 1	117 910 1	111 5/11	29.759	51.358	87.480	90.640	100.150
	NT2RP2000426	114.626	117.810	111.501					
	NT2RP2000428	56.117	63,709	38. 237	12.835	20. 360	38. 761	41.161	42.507
5	NT2RP2000438	54.621	34.534	49.392	15.765	13.700	27. 527	31.816	22.667
3	NT2RP2000447	41.157	17.807	23.084	6.863	12.794	25. 289	17.738	11.474
	NT2RP2000448	26.410	27,807	28. 584	7.787	12.459	20.751	18. 164	12.208
	NT2RP2000459	44. 499	36.093	89.605	12.882	14. 284	17.465	15. 331	9.860
	NT2RP2000479	21. 922	30.183	53.808	9. 553	8.835	9, 648	8.854	6.739
				207.697	30. 335	41.292	29.900	25.090	43.440
	NT2RP2000498	97. 221	94.691			10. 196	9, 766	10.763	12.056
10	NT2RP2000503	15.067	15, 551	20.810	5. 166				
	NT2RP2000510	8. 340	5, 361	8.647	4, 438	7.160	4. 784	7.812	3, 890
	NT2RP2000514	10.423	8.148	14.693	2.596	1.773	12.792	6.695	2.902
	NT2RP2000516	24. 587	13.672	21.344	7.854	6. 333	13.895	7.395	10, 960
	NT2RP2000523	10.281	2, 981	4. 878	1, 371	8. 071	0.000	6.857	1.961
	NT2RP2000533	26. 452	20, 054	30, 481	4. 391	7. 528	16.125	48, 840	17. 396
	NT2RP2000540	52. 523	22. 512	28.503	13.567	14. 512	28. 427	22. 545	11.372
15		22.542	17.741	11.176	7, 337	26.779	12.216	8. 288	6.918
	NT2RP2000547					30.047	43, 813	31, 490	9. 367
	NT2RP2000557	91.024	63, 951	163.497	30, 438				
	NT2RP2000558	53.959	47.359	125.971	27. 348	16.844	24. 191	17.114	21.905
	NT2RP2000564	30.445	23.046,	22.258	13.084	14. 165	16. 265	14.861	11. 150
	NT2RP2000565	12. 593	5.857	10.293	5.077	0.000	4.189	5, 009	9.707
	NT2RP2000583	92. 921	56.070	68.992	29. 211	14. 291	50. 282	32.844	34.467
20	NT2RP2000591	14.655	9, 331	13.087	3.504	0.000	10. 526	4.362	2.073
	NT2RP2000599	8.002	4,780	- 7. 951	1.807	1.614	6.232	2.299	8.293
	NT2RP2000601	63.609	21.655	47. 106	9.673	13. 430	48. 855	32. 575	8.428
	MT2RP2000603	101.578	37.142	48. 248	16, 412	25. 194	51, 543	39. 363	20. 157
				110.636	42.146	27.855	28. 332	30.624	31.736
	NT2RP2000610	78.342	66,011				83.079		185, 276
	NT2RP2000614	139.380	106.590	188.604	171.750	58. 678		86. 298	
25	NT2RP2000616	124. 143	34.073	58.053	15.031	27. 800	81.174	49.504	27.143
	NT2RP2000617	50.724	37.802	37.086	17, 602	12.086	34, 751	20. 157	16.389
	NT2RP2000623	39. 247	19.740	34. 797	9.070	10. 223	19.775	10. 261	13. 251
	NT2RP2000634	29.431	24.224	35.865	13.077	19.480	16.373	23.806	11.338
	MT2RP2000636	39, 598	28.832	34. 563	11.868	13. 914	14, 342	6.334	10.485
	NT2RP2000638	43. 027	34, 379	58. 259	14.094	15. 200	22.724	21.525	6.843
••	NT2RP2000644	87.622	66.336	227. 352	37.298	35.466	29. 256	23.666	11.793
30	NT2RP2000649	28.849	24.035	32.562	15, 166	18.629	25.012	15, 485	15, 528
		39. 595	25.065	30. 965	10.579	14. 587	24. 849	13. 667	10.824
	NT2RP2000652			7. 925	2.952	4. 388	9, 997	3. 990	6.959
	NT2RP2000656	12.851	14.986					3.050	2.703
	NT2RP2000658	8. 192	5.499	7.563	1.162	3. 535	5. 669		
	NT2RP2000663	38.633	21.653	37.840	5.964	12. 174	20.777	13. 553	39.917
35	NT2RP2000664	102.627	41.981	90.611	25, 300	30.038	73.440	66. 686	30. 192
33	NT2RP2000668	41.209	35. 434	46.568	16.251	14. 705	25, 339	29, 016	11.020
	NT2RP2000678	6.908	2.096	21.949	0.402	5. 899	0. 262	1.098	1.488
	NT2RP2000694	47.376	19,985	45.832	2.636	16.192	24. 523	19.843	12.311
	NT2RP2000704	159.158	114.202	205.746	44.471	48.627	68, 161	47. 919	40.349
	NT2RP2000710	33.138	26.994	21.890	10.683	6.833	17. 938	13. 596	8.070
	NT2RP2000712	15.016	11,689	29.736	12.471	8.668	17.629	19, 970	23.796
40	NT2RP2000715	61.771	35, 912	115.757	20.470	17. 051	26, 042	17. 159	21. 325
-	NT2RP2000720	38. 951	26.992	43.620	14.647	11.930	21.500	23. 895	26.128
							2, 127	4. 657	5. 827
	NT2RP2000731	8.039	11.373	11.261	2.986	4. 755			
	NT2RP2000739	83.662	28.893	61.699	15.623	21.878	30,716	28. 485	17, 190
	NT2RP2000748	21.953	22.377	38. 996	16.815	15. 564	15.846	20. 219	20.054
	NT2RP2000749	46.622	49.334	65. 231	13.317	57. 514	52.159	26. 941	23.858
45	NY2RP2000758	79.204	43.258	49. 581	18.768	17.058	49. 245	31.463	9. 472
	NT2RP2000764	65.396	28, 914	41.243	10.203	15, 308	36.761	32. 438	13, 134
	NT2RP2000766	40.275	50,060	83. 340	10.610	48. 180	26.508	18.850	15.663
	NT2RP2000777	92.029	39, 471	41.396	32.309	33.513	94.887	43.480	40.212
	NT2RP2000786		61.265		16.798	30.669	51.517	41.968	37.840
		91.676		70. 189					
	NT2RP2000793	245. 992	91, 135	151. 153	57. 903	62.361	191.087	132.793	68. 352
50	NT2RP2000796	24.053	16.664	26. 593	12.773	8. 297	14.258	11.004	10.440
	NT2RP2000809	118,982	88.958	221.024	42.198	50. 535	65.921	39, 243	46. 532
	NT2RP2000812	23. 931	28.037	26.224	15.476	9. 968	23. 492	19.671	6.489
	NT2RP2000814	9. 108	7.645	7.698	5. 179	4. 196	5. 655	3. 821	2. 231
	NT2RP2000816	49, 615	22.174	23.358	8.758	4. 975	23.109	16.789	17. 124
	NT2RP2000818	8. 156	2.591	1.260		0.840	1,656	0. 942	0.250
	MI ZN. 2000010	3. 130		1.200	1 3.752	1 4.0.4	1		1

492

Table 83

	NT2RP2000819	18.931	14.180	22. 186	4. 470	4. 973	11.654	8. 535	5. 164
	NT2RP2000841	28.455	24.097	27. 497	8. 335	10.021	20.722	20. 951	21.582
		34. 381	17.071	34.845	8.688	13.092	22.498	16. 807	14. 291
	NT2RP2000842				60, 191	54, 194	71.809	58, 470	61.375
1	NT2RP2000845		19. 456	21, 479	5. 334	8.450	25. 326	17.757	8. 334
	NT2RP2000863	43.408	45, 920		29, 897	15. 173	32.007	24.723	27.411
	NT2RP2000880	57.370	13.581	18. 264	3, 215	4. 302	10.360	11. 152	10. 295
	NT2RP2000892	10.063	18, 305	26. 241	9. 579	7. 189	24.935	24. 193	12.407
	NT2RP2000894	64. 414	14. 595	23.755	3. 839	8.850	15.625	14. 467	11.189
	NT2RP2000903	38. 945	24, 347	34. 459	12. 388	12.400	29.304	18.688	23. 184
	NT2RP2000906	43.895	47, 430	175. 193	28. 258	21.020	30.976	28.638	37. 229
	NT2RP2000910	76.036	104. 907	108.794	52. 697	65.250	51.718	29. 344	55, 383
	NT2RP2000931	68.351 30.706	39. 023	31.030	6. 448	13.290	15. 553	11.313	11, 145
	MT2RP2000932	55, 079	37, 641	47.798	12.045	19.899	32.600	18.046	21.528
	NT2RP2000938	54.610	32.689	54, 181	11.802	18.241	33.817	55. 424	23. 572
	NT2RP2000943	20. 426	12. 332	17. 780	3, 161	5. 343	6.479	8.015	4. 052
	NT2RP2000957	74.825	23. 934	37.910	10, 227	22.164	41.633	29. 369	21. 256
	NT2RP2000959	15.840	25.061	17. 980	5. 521	4.208	9.176	3. 539	6.349
	NT2RP2000965	52.687	40. 458	51, 330	27.882	16. 372	29.535	32.993	35. 543
	NT2RP2000970	84.866	72.715	196. 279	29. 249	36.529	42.914	24.489	33. 313
	NT2RP2000973	42.690	30, 786	42. 102	8.964	13.498	23.369	20.702	18.360
	NT2RP2000985	33. 281	22.399	26.930	8.628	4.869	20.022	22. 445	14.030
	NT2RP2000987	47.735	66. 487	94.477	25.911	19.844	27.890	23.633	33. 551
	NT2RP2000997	42, 801	43, 070	56.966	15.270	16, 292	49.613	53.625	99. 622
	HT2RP2001024	47, 605	28.976	34.658	13.810	14.526	32.054	39. 269	22. 962
	NT2RP2001028	32.502	24,770	88.599	12.437	11: 259	13. 181	13. 919	9. 824
	NT2RP2001036	206. 163	234. 625	568. 339	116.746	85.893	125.996	88.623	100.568
	NT2RP2001039	26.909	37.527	31.356	6.335_	15, 429	17.827	107.341	12.412
	NT2RP2001044	51.134	33.868	42.988	9.015	23.633	31.422	25.682	20. 463
	NT2RP2001056	84.875	95.778	164. 256	33. 325	35.039	45.764	30.831	44. 980
	NY2RP2001065	57.092	61.052	49. 599	18.558	20. 229	29.013	30.628	32. 966
	NT2RP2001067	17.223	18.596	14.258	5. 284	6.021	4. 582	10.045	8. 782
	NT2RP2001070	92.615	68.975	230.584	37.646	41, 225	36. 295	43. 293	26. 959 45. 105
	NT2RP2001081	134.654	80.124	269.700	35. 425	37.697	42.849	35.852	15. 956
	NT2RP2001087	54.476	40.059	74.079	12.377	21.043	25.654	22.663 4.258	5. 446
	NT2RP2001094	11.558	8.400	11.506	4, 416	3.583	4.503	41. 672	39, 968
	NT2RP2001119	66.924	57.741	177. 347	36.523	37. 388	40.013 16.757	28.865	13. 483
	NT2RP2001127	52. 585	39, 380	36. 247	9.959	18.625	28.836	28. 731	38. 218
	NT2RP2001133	94, 638	97.465	155. 477	25. 417	36. 346	20. 107	24.686	30. 341
	HT2RP2001137	61.770	53.486	51.726	12.991	40.072 14.588	17. 434	23.807	20.602
	NT2RP2001142	54. 131	38.507	34. 342	8. 552 17. 462	14.077	23.064	30. 676	22.427
	NT2RP2001149	96.617	49.914	71.348	65. 294	77.914	146.883	169. 121	159. 484
	NT2RP2001168	313.055	217.008	22.710	16.143	12.538	13.238	14. 902	9. 473
	NT2RP2001173	25. 149	27. 272 17. 440	22.879	11.089	14, 190	18. 125	50.600	22.839
	NT2RP2001174	21.134 99.803	60. 549	84. 254	29.471	35. 438	70. 558	65.859	57. 928
	NT2RP2001184	19. 492	14, 580	26.749	5. 551	9.060	20.695	9.289	15. 340
	NT2RP2001200	39, 331	44. 223	52.647	14.745	26.231	26.146	33. 102	31.874
	NT2RP2001218	32. 396	16.531	28, 960	21.387	13.855	8.618	18.872	11. 236
	NT2RP2001223	86. 393	27. 183	46.400	14. 290	23.545	53. 375	28.096	26.084
	NT2RP2001226	223.868	143,880	155.700	46. 575	60.808	148.876	100.150	92.898
	NT2RP2001227	100.969	51.807	65.094	19, 398	24. 302	57.877	36.375	36.204
	NT2RP2001232	49.733	30. 526		11.691	29.542	27.238	22.294	35. 950
	NT2RP2001233	42.734	36.288		58. 935	18, 921	38.027	28. 582	69. 539
	NT2RP2001245	28, 251	16.266		18.419	8.746	38.272	8.565	38.035
	NT2RP2001246	24.708	44.426		19.345		35.994	31.550	37.123
	NT2RP2001268	44. 328					54.014	31.715	54.645
	MT2RP2001270	37.478		29,740			24.740	26.469	21.423
	NT2RP2001276	15. 931	7.906	12.674			12.549	17.273	10. 235
	NT2RP2001277	22.937					12.672	2.878	16. 107
	NT2RP2001290	66.867	20.688				38.444	27.073	34. 029
	NT 2RP2001295	22.717	21.639				24. 206	7.863	10. 592
	NT2RP2001297	105.753					152.615	178.985	
	NT2RP2001301	47.099	37.782	53.504	25. 117	15. 392	49. 389	38.668	29. 281

#### .. Table 84

	NT2RP2001312	493.097	175. 989	324. 513	96.070	132, 150	315.768	282. 270	146.542
		188, 839	50. 032	95.732	33.162	58.029	112,666	87.335	71,442
	NT2RP2001327	177, 255	162. 267	495. 438	96.591	104. 203	93. 675	57.120	68.709
5	NT2RP2001328		92. 245	40. 237	32.288	34.069	91. 368	77. 221	45. 741
	NT2RP2001341	196.358			72.828	67.867	81.012	36. 464	72, 260
	NT2RP2001347	148. 143	157. 594	486.643		96. 521	146.562	77.918	108.669
	NT2RP2001366	160.323	170.553	496.412	116.205 31.128	51.690	147, 191	118. 132	56.442
	NT2RP2001378	217.791	51. 524	110.978			8. 483	2.703	10.418
	NT2RP2001381	16.578	13. 963	19.068	15. 119	9. 576	52. 881	33. 168	44, 592
10	NT2RP2001388	84.013	52.476	228.213	47.276	49.007			127.811
	NT2RP2001391	806.136	1438. 949		960. 225			23. 261	
	NT2RP2001392	56.943	65. 258	70.204	19. 962	26.883	46. 456		14.231
	NT2RP2001394	104.258	120.852	350.764	78.963	59.635	75. 686	42.505	53.751
	NT2RP2001397	37.759	22.378	38.780	40.524	15. 364	21.089	16.393	16.560
	NT2RP2001400	24. 214	10.586	19.685	10.414	12. 173	24. 380	11.798	22.055
45	NT2RP2001408	34. 405	28. 262	69.823	33.071	21.313	29. 278	20.555	45.713
15	NT2RP2001420	74.700	70.462	212.932	44, 495	49.469	33. 427	30.009	41.019
	NT2RP2001423	20.045	17. 202	38.815	16, 204	11.082	21.739	12.751	10. 452
	NT2RP2001427	88.620	91.272	206.946	51.057	36.829	49.854	34. 587	57.012
	NT2RP2001428	47.617	45, 465	55.112	19, 580	15. 421	24.651	10, 915	29. 985
	NT2RP2001436	19.654	25.608	50. 345	11. 202	18. 548	32.033	22.720	5.351
	NT2RP2001440	11.871	12. 123	19.145	7.724	5.414	7.413	19.955	16. 145
20	NT2RP2001445	11.934	7.217	22.053	5.885	6.872	7.794	3.377	20.818
	NT2RP2001449	20. 271	20. 423	53. 385	13. 242	8.026	-8. 439	6.342	9.186
	NT2RP2001450	47.497	32.496	58. 237	18.660	21.208	28.880	23.620	30. 207
	NY2RP2001467	40.279	40.050	115.089	25. 502	21.744	18.716	21.445	37.772
	NT2RP2001469	66.890	35.784	93.465	23. 588	33. 470	54.095	54. 103	33.386
	NT2RP2001480	69.698	53.669	54, 777	16.208	26.373	44, 943	30.622	26.208
25	NT2RP2001495	14.156	12.199	18.018	8. 178	14.762	10.694	8.800	14.613
	NT2RP2001499	40.983	50. 266	57.334	23. 302	22.298	37.271	26.788	35. 187
	NT2RP2001506	83.528	66.377	104. 162	41.795	65.692	61.567	55.661	35.667
	NT2RP2001508	25.746	36.879	44.112	33.620	14.149	23.999	19.783	36.174
	NT2RP2001511	231.898	147, 751	199.611	46. 927	77. 381	122.787	130.829	108.021
	MT2RP2001514	121.671	47. 391	103.398	24.149	31.957	72.965	63.365	38.173
30	NT2RP2001520	38.773	20.470	34.140	14.159	13. 366	19.602	22.077	7.741
•	HT2RP2001526	102.469	96.418	139, 331	62.159	83. 922	85.309	60.450	66.763
	NT2RP2001529	189.308	69.082	103.704	31.713	54. 543	173.158	96.700	74.482
	MT2RP2001536	22.047	14. 186	19.269	9. 553	7.196	16.531	13.646	17.343
	NT2RP2001538	123. 315	222.563	281.173	191.775	90. 257	199. 255	133.592	422. 435
	NT2RP2001547	45, 201	33.999	42.028	12.917	14.746	31.438	29.406	24.085
35	NY2RP2001560	146.079	68.501	131,623	35.625	46.061	88.704	90.584	78.703
33	NT2RP2001562	53. 975	35.141	47. 262	23. 297	18.361	43.041	30.635	47.577
	NT2RP2001566	55. 453	48, 563	91.463	37. 157	27. 507	54.780	37. 595	42.663
	NT2RP2001569	131.940	142.523	361.640	62.136	60.136	90.021	46. 500	62.567
	NT2RP2001576	103.537	76.306	58.434	23.607	34.646	91.308	67.270	45. 219
	NT2RP2001581	149.528	208. 681	239. 575	139, 522	72.883	196.577	126. 583	231.505 43.012
	NT2RP2001597	52.409	27.790	43.630	13.807	18.650	35.875 22.525	13.760	29.828
40	NT2RP2001601	33.796	37.430	70. 562	17.535	15. 251	9.119	9.830	14.501
	NT2RP2001613	10.438	5. 350	6.715	3. 155	6, 423	50.117	31.175	117.652
	NT2RP2001628	87.399	43, 401	48.713	17.774	25. 577	30.530	21.235	42.849
	NY2RP2001634	38.792	56.546	47.793	23.992	16.006 36.011	40.036	38.853	22.210
	NT2RP2001635	63.818	69.842	156. 279	31.411	11.137	20.048	20.365	48, 159
	NT2RP2001660	31.664	25.538	25. 905	6.081 52.514	43.761	63.759	56.518	43.557
45	NT2RP2001662	122.557	88.914	242. 932		16.477	39. 485	20.869	25.608
	NT2RP2001663	33.056	34.206	58.783		26, 225	33.037	25.666	35. 948
	NT2RP2001672	51.656		140, 882		5. 806	3. 149	4.861	6. 461
	NT2RP2001675	8. 589	6. 791				40.323	46.465	47.741
	NT2RP2001677	51.810	49, 851	68.423		27.233 46.915	65.668	62.835	64.186
	NT2RP2001678	70.100				10.015	9. 959	25. 390	9. 277
50	NT2RP2001683	16.088	14.728				64. 457	41.391	50. 672
	NT2RP2001699	116.996	54.743			34. 234	58.720	45.599	68.302
	NT2RP2001707	94.748	66.728 33.745				38.973	31.931	30.227
	NT2RP2001720						69.464	66. 522	35.468
	NT2RP2001721	73.164 23.081	35.354				21.055		28.546
	NT2RP2001740	23.081	30.430	27.131	1 12. 949	12.240	1 41.033	1 20.033	1 20.040

494

Table 85

	NY2RP2001748	164, 370	51.538	151.756	22.608	36.134	86.312	52.323	35. 004
	100 200 000 100	10, 363	5.707	7. 354	3.303	2.490	20. 122	2.133	9. 371
	NT2RP2001755	10.743	10.704	7.130	4.777	5. 548	16.360	7.429	3.763
5	NT2RP2001762	122.047		129.000	29.098	38.722	67. 999	58. 129	48.111
_	NT2RP2001768	29. 307	28.706	32.455	11.608	15.175	19.399	20.505	29. 469
	NT2RP2001769	18.824	19, 322	24. 434	8. 167	13.814	14.835	14. 266	10. 332
	NT2RP2001784	111,510	63.886	82. 038	33, 170	41.704	47, 921	62.218	54. 508
	NT2RP2001805		10. 225	13.797	4. 221	9.786	3,548	11.805	8, 246
	NT2RP2001813	15.000	12.403	19. 383	6.848	8. 320	6.884	10.608	15. 163
10	NT2RP2001817	30. 494	21.374	23.441	6.808	14. 438	12.858	13.055	9. 397
	NT2RP2001818	153. 478	143. 980	348. 522	65.249	56.344	69.434	48.042	62.813
	NT2RP2001837	68, 237	44.006	65, 237	21.186	23.824	37.874	35. 524	54. 235
	NT2RP2001839	45. 604	33.558	72.763	21.180	25. 185	40.479	31.542	29. 326
	NT2RP2001861 NT2RP2001869	79, 101	52.967	123. 399	29.766	25, 811	40.870	38. 251	38. 150
	NT2RP2001876	20.847	28, 536	35, 991	18.044	13, 257	29. 195	20.056	35.651
15	NT2RP2001878	105. 429	34.989	86.887	21.675	33.547	76.806	64.301	35. 521
	NT2RP2001881	25. 562	5, 186	16, 935	8. 594	6.002	8.017	5. 474	16.018
	NT2RP2001883	152. 487	96. 494	76.800	26.663	40.257	93.069	57.806	50.662
	NT2RP2001884	40.027	29. 435	18, 175	19.127	0.000	34.665	13.313	27. 989
	NT2RP2001885	41.527	29. 494	60.284	13.719	9, 345	26. 427	24.717	30.448
	NT2RP2001898	152.071	65.585	135.420	33.617	41.173	112.042	64. 105	57.703
20	NT2RP2001900	20.075	16. 336	54. 207	10.431	9.790	20.098	19. 168	30. 123
	NT2RP2001903	389. 922	207.168	314.475	131.627	170.618	361.733	261.185	289. 339
	NT2RP2001907	118.240	77.557	213.664	50.816	46.691	58.895	52.711	56.061 15.535
	NT2RP2001915	29. 335	9. 240	29. 213	5. 804	10.101	8.718	14.671	30.244
	NT2RP2001921	70.657	42. 199	23. 786	27.411	23.817	52.083	27.655 37.077	26, 703
	NT2RP2001926	86.771	11.953	10.434	11. 123	10.945	27.144 114.539	90. 251	48. 849
25	NT2RP2001933	210.457	80.003	159.875	38. 312 9. 560	53. 192 6. 311	8,705	3. 968	4. 244
	NT2RP2001936	9. 271	13.789	9.841	96.135	99. 997	227.342	186.800	161.131
	NT2RP2001943	329.800	151.136	357.167	18.830	11.786	20.082	32.636	29. 552
	NT2RP2001946	35.700	27. 839 40. 322	58.250	17. 399	25. 524	30, 411	31, 309	15.258
	NT2RP2001947	49. 825 6. 858	5, 149	39.338	5. 855	16.449	8.590	3. 943	39. 227
	NT2RP2001948 NT2RP2001956	204. 499	97.036	150. 184	34, 215	55.776	144.746	109.645	45.142
30	NT2RP2001969	63.044	42.091	64.895	18.446	22.555	64. 128	29.876	27.818
	NT2RP2001976	8.014	10. 925	13. 322	14. 259	2.776	2.729	6.432	21, 452
	NT2RP2001978	60.910	40. 459	87.051	23. 282	28.689	25.497	33. 528	35. 507
	NT2RP2001985	73.126	35.661	72.052	21.029	30.385	52. 486	46.885	41.899
	NT2RP2001991	32.897	34.028	33, 239	10.548	15. 586	20.531	18. 489	33. 157
	NT2RP2001997	38. 265	33.006	69.711	20.057	29.835	29.074	30.213	39.156
35	NT2RP2002015	341.650	572.382	454. 288	330.114	80. 297	366.270	346. 254	476.966
	NT2RP2002017	33.458	25.736	55. 897	13. 982	18. 424	23.720	92.718	17.897 55.437
	NT2RP2002025	201.899	111. 493	125, 922	38.775	57.018	118.130	65.007	88. 254
	NT2RP2002030	147.806	150.643	447. 960	95.773	58.859	95, 260 127, 664	86.380	56.817
	NT2RP2002032	170. 695	55. 335	101.868	30. 495 84. 872	61.493	72.667	37. 144	74. 278
40	NT2RP2002033	147, 111	92.379	481.152 17.284	5.762	7.552	5. 398	10.885	30. 538
40	NT2RP2002041	15.097	12.379	25, 228	11.030	8.158	11.642	14, 255	15. 385
	NT2RP2002046 NT2RP2002047	19. 261	15. 499	12.076	6.530	14. 384	9.918	10. 225	22. 164
	NT2RP2002050	71. 226	75, 633	97.017	33.238	36.421	49.003	42.580	45.656
	NT2RP2002052	75. 004	67. 588	69,616	25, 123	25.691	49.820	32.819	35. 546
	NT2RP2002058	9.803	11.955	11.648	6. 527	5.940	8.570	15.678	16.434
45	NT2RP2002060	147, 927	40. 191	79, 254	17.661	30.022	83.968	55. 933	35.933
13	NT2RP2002063	8. 334	10.615	17, 124	3.910	9.032	6. 499	6.095	43.967
	NT2RP2002066	85, 296	31.968	71.727	16. 597	28. 928	52.589	40.814	37. 383
	NT2RP2002070	24.791	21.309			11.537	11.893	11, 300	28.065
	NT2RP2002076	28. 441	16.541	17.729			13.321	11.910	9. 273
	NT2RP2002078	75, 992					65.434	39. 358	28. 599
50	MT2RP2002079	15. 378	6.595				7, 129		27. 362
	NT2RP2002099	78. 520					51.830		
	NT2RP2002105	45. 619					33.970		26. 203
	NT2RP2002115	4. 270							
	MT2RP2002124	9. 528							
	NT2RP2002137	42.205	16.239	58.339	7. 326	12. 132	22.097	1 14. 004	1 .5.003

Table 86

									** ***
	NT2RP2002139	134.906	45. 566	87.430	23.245	34.053	84. 389	66.892	39. 692
	NT2RP2002154	73, 877	40. 956	58.084	17.359	21.276	53.977	32.847	25, 981
		312.813	448, 404		246.578	165, 102	220, 200	117.089	396.951
5			30. 674.	55.659	15, 991	11.479	50.028	14. 567	76.933
	NT2RP2002172	30. 233		31.771	9. 436	13.476	23, 455	23.637	18.868
	NT2RP2002185	35. 127	22.047			58.463	164, 587	118.094	78.046
	NT2RP2002188	281.595	70. 032	141.729	44. 333				25.727
	NT2RP2002192	28.830	19. 474	84.849	18.536	13.421	8. 267	10. 908	
	NT2RP2002193	51.545	23. 270	33.672	10.534	17.989	33.897	31.972	33.050
	NT2RP2002208	28. 592	23. 922	46. 525	15.986	13.078	25. 948	18.689	40. 263
10		13. 529	18, 299	23.304	8.697	7,005	20, 832	6. 994	4. 421
	NT2RP2002219			18.238	6. 451	5.394	5. 290	2.444	1.640
	NT2RP2002231	3. 623	9, 145			12.646	31,637	16.070	23, 193
	NT2RP2002232	41.922	30,600	40.665	10.290			12. 594	16.053
	NY2RP2002235	25. 174	12.829	11.461	1.747	8.624	10. 246		
	NT2RP2002239	123.883	99. 627	183.537	54. 220	35.311	68.845	72. 486	114. 538
	NT2RP2002252	173. 209	45.051	80. 502	16.296	33.546	82. 843	82.445	52.048
15	NT2RP2002256	6.776	3, 892	.12.301	3.488	7.236	6.566	9. 391	- 9.526
		14.914	18.059	11, 330	3.304	7.442	11, 747	12.965	135.057
	NT2RP2002257			41. 590	9.164	7.968	18.892	22.893	29.020
	NT2RP2002259	25.623	20. 902			7.884	26.448	8.234	20.246
	NT2RP2002264	35.467	21.3801	27. 456	3.962		63.895	30.469	55. 401
	NT2RP2002267	99.224	90. 968	353.970	55.091	43. 373			19, 108
	NT2RP2002270	12.038	20.146	13.141	7.551	3.523	7.777	6. 701	
20	NT2RP2002281	49.615	38.410	43. 936	21.926	17.935	51. 455	14.825	35. 239
	NT2RP2002288	18.840	15. 310	15. 237	4.623	6.951	4. 505	6.438	4. 321
	NT2RP2002292	70.138	79, 487	98.062	32, 152	32.815	48.306	41.287	55. 682
		28. 411	21, 790	28. 450	15.762	10.016	23.812	12.394	31.923
	NT2RP2002299		27. 505	25. 401	9.478	10.570	14.112	10.173	10.213
	NT2RP2002304	17.776				10.392	41, 845	21.011	16.815
	NT2RP2002312	32.053	25.004	19.733	5.118			17.862	43.519
25	NT2RP2002316	15.618	29. 406	20.363	11.321	29.588	16.866		
	NT2RP2002325	32.321	23.882	28.697	6.692	9.875	26. 435	21.261	36.989
	NT2RP2002333	117.384	75.765	92.724	37.475	55. 245	56.768	79.089	134.509
	NT2RP2002371	35.025	49. 789	54.117	20.073	31.179	10.486	24. 281	48.279
	NT2RP2002373	73.024	55. 638	58.797	24.729	33.686	48.754	58. 440	58. 483
		4.610	6.610	5. 950	2.906	4.109	10.398	7.035	3.142
	NT2RP2002381			39, 973	10. 268	23.738	57.377	29.062	18. 367
30	NT2RP2002385	73.600	28. 798		1.941	3.227	11. 225	3, 017	2.611
	NT2RP2002394	4.749	3.341	5. 573			22.778	22. 751	14.463
	NT2RP2002408	30. 199	16.610	24.803	8.840	17.966			
	NT2RP2002409	466.226	415. 995	746.844	183.086	221.410	247.550	215. 812	235.852
	NT2RP2002424	73.955	40.022	38.701	11.417	27.269	38.757	35. 192	25.977
	NT2RP2002426	42.246	46.209	138.641	18.951	43.167	21.993	14. 146	29.925
05	NT2RP2002429	38.798	37.515	37, 290	13.976	31,959	40. 592	16. 576	28.408
35	NT2RP2002437	41.182	44.109	103.486	16.002	6.706	22, 769	11.006	18.502
			110.081	147.018		60.331	171.025	155. 332	90, 923
	NT2RP2002439	300.787				21.412	43, 427	38. 136	78.512
	NT2RP2002442	51.674	59. 162	57. 683			42.083	43.069	38.125
	NT2RP2002457	87.804	91.782			50.903			34.847
	NT2RP2002464	97.665	38.612		20.743	31.183	66.794	48.779	
40	NT2RP2002475	87.229	49, 226			38.579	51. 432	45. 816	27.604
70	NT2RP2002479	43. 495	20. 134	24. 184		13, 868	35, 366	19. 292	22.684
	NT2RP2002487	95.041	44. 922	72.897	21.815	31.046	43.590	37. 943	47.177
	NT2RP2002498	32.022	15.599			8.092	15. 582	24. 301	15, 152
	NY2RP2002503	143.137	80. 337			35.509	96.570	63.743	69. 363
			12.130				16.936	24, 798	15.731
	NT2RP2002504	28.779					192.079	171.751	115.972
45	NT2RP2002510	389. 826	185.539		1 11 000		38. 504	25. 659	37.802
	NT2RP2002520	28.465	20.629			14.948			45. 539
	NT2RP2002527	82.404							
	NT2RP2002533	453. 205							188.717
	NY2RP2002537	39.475		89.50	4 25, 635				23. 210
	NT2RP2002542	68.000				29.448	38.068	33.806	
	NT2RP2002546	27.656						6. 935	5. 27
50									
50	NT2RP2002549	41.394							
	NT2RP2002564	135.808							
	NT2RP2002591	34.917							
	NT2RP2002595	29.155	28.99						
	NT2RP2002602	62.164	42.49						
	NT2RP2002606	23.368		1 18.05	8 7.40	14. 39	5.066	8. 402	33.19
55									
· <del>-</del>									

Table 87

						Y			44 405
	NT 2RP2002609	51.566	22.622	50.513	17.534	20. 249	18.692	26.812	44. 491
	NT 2RP2002618	54. 802	20. 530	64.541	20.957	20. 177	31. 222	20.834	32.080
				361.642	75.866	73.104	87. 556	37.662	72.940
5	NT2RP2002621	108.854	151.631						
3	NT2RP2002643	79.459	49.749	159. 126	32.285	31.588	30.054	50.389	48. 139
	NT2RP2002672	97.309	70.875	124.816	41.317	54.912	65. 362	54. 912	61.191
			27. 367	31.454	11.741	15. 225	18. 592	18.872	41.668
	NT2RP2002673	33. 731							
	NT2RP2002674	13.503	12.059	23.980	5. 008	15.903	5. 926	8.720	8.883
	NT2RP2002686	45, 156	22.604	57.057	22.253	22.373	30.389	27.672	13. 377
		85. 273	71, 163	154.737	61.783	35.115	56. 421	42.460	68.118
10	NT2RP2002588						43, 453	30.540	28.172
10	NT2RP2002695	80.865	40.613	62.941	15.213	22.197			
	NT2RP2002701	68.274	58.034	54.220	24.008	29.811	75. 585	54.744	29.997
	NT2RP2002706	66,710	49.408	147.083	42, 409	25.501	40.462	31.482	31.678
						312.053	990.051	876.290	401.334
	NT2RP2002710	876.030	389.806	785.892	246.642				
	NT2RP2002721	120. 344	48.897	112.902	26.906	37.076	81.599	62.600	40.801
	NT2RP2002727	19.985	16.809	28.658	5.885	10.968	18. 932	17.127	19. 197
15		84, 484		244.997	57.973	45.229	35. 711	33.199	39.655
13	NT2RP2002734		81.389						
	NT2RP2002736	18.170	7.757	29.873	5. 264	10.456	10. 179	9. 257	11.010
	NT2RP2002740	13.219	14.424	23.343	12,863	6.975	8. 152	8.795	7.772
				223. 592	33.955	36.594	51.261	45. 295	14.049
	NT2RP2002741	77.823	67.266,						72.280
	NT2RP2002750	140. 558	111.369	512.500	99.357	68.412	72.711	76.999	
	NT2RP2002752	177.349	105. 312	290.520	63.592	64.508	103. 376	92. 228	65.849
20	NT2RP2002753	131.824	60,851	110.980	32.981	43.667	85, 850	102.908	117.429
20						37.588	59. 420	51. 267	51.768
	NT2RP2002760	130.675	58. 967-	119.405	28.837				
	NT2RP2002769	19.077	14.018	32.873	14.190	12. 132	10. 357	15.988	25.043
	NT2RP2002778	38,616	37.548	30.303	18, 271	16.022	71.865	31.460	77.045
			55. 458	105.096	34.190	38.076	66, 995	54.639	45.519
	NT2RP2002791	95. 319							
	NT2RP2002800	90.052	59, 554	197.798	40, 413	37. 123	87.119	52.880	48. 173
25	MT2RP2002805	14.997	12,041	9.573	4.470	8.397	5. 324	5.699	14.665
25	NT2RP2002811	84. 563	36.955	70.308	17.273	24, 509	89,018	46.163	49.186
							56.385		38.118
	NT2RP2002824	44. 392	48. 364	75.269	21.980	25. 621		42.073	
	NT2RP2002839	45.683	28.499	42.893	12.083	18. 567	22.078	23.650	21.604
	NT2RP2002845	46.337	22.545	45.003	11.450	16.060	6.978	26, 900	14.552
		26.773		27.648	7.358	7.968	15, 413	17.314	11.937
	NT2RP2002857		11.114						50, 334
30	NT2RP2002862	122.430	114.903	392.000	81.893	61.001	82.758	60.301	
30	NT2RP2002880	46.913	32.677	29.822	12,750	16.704	35, 359	14.768	24.866
	NT2RP2002885	24. 335	26.185	27.174	10.146	19,062	54. 580	55, 170	22.593
							38.658	34. 150	26.201
	NT2RP2002891	33.411	27.772	38.018	14, 500	16.632			
	NT2RP2002907	31.117	36.465	35. 948	13.227	13.010	49. 135	37. 225	26.747
	NT2RP2002925	30.213	17.281	33, 298	11.072	11.726	25. 559	24.754	17. 499
			35.383	40.539	21.437	7.365	35, 485	14.771	39.460
35	NT2RP2002927	21.224							
55	NT2RP2002928	13.771	14.521	49.574	11.977	6.869	9.129	7.289	8.057
	NT2RP2002929	21.741	22.530	32.027	7.934	12.601	20. 143	13.573	25. 568
	NT2RP2002934	63.248	35. 331	42.688	10,849	16.987	39, 637	27.937	23.467
					15, 330	19.313	35. 512	35.749	26.290
	NT2RP2002939	53.914	30.833	62.082					
	NT2RP2002942	82.129	82.694	187.805	50. 572	53.315	49.000	38. 922	90. 399
	HT2RP2002954	33.490	25, 335	35.779	11.591	11.217	27. 293	16.672	26.618
40	NT2RP2002959	18.029	22. 305	18.230	8.391	14.540	12.392	9. 227	31. 203
40						18. 382	28.562	36.888	41.144
	NT2RP2002974	34.775	17.807	29.755	6, 220				
	NT2RP2002976	7. 266	6.893	13.152	2.886	5. 205	17.007	6.657	17.861
	NT2RP2002979	156, 906	139, 229	395.529	82.939	71. 144	104. 220	76.074	81.377
				285. 396	49, 557	40.675	57.510	33.004	50.480
	MT2RP2002980	98.467	79, 422						
	NT2RP2002986	210.452	66.962	105.842	25. 570	34.404	156.863	99.482	35.944
45	NT2RP2002987	170, 131	130,848	355. 987	1114,067	85.014	125. 562	105. 241	119.400
10	NT2RP2002988	35.092	33.804	42.437	7.516	22.093	78.216	26.257	53. 462
								1 2 22	1 . 4 44.
	NT2RP2002993	41.408	20.150	29.978	8.083	13. 951	19.869	17.068	17.776
	NT2RP2003000	91.683	72.701	265.303	52.674	45. 922	52. 225	38. 486	61.960
	NT2RP2003008	19. 429	42, 300			11.323	22.796	23. 430	31.344
						198. 298	95, 120	89. 298	74. 362
	NT2RP2003020	146.283	83.102		31.287				
50	NT2RP2003032	42.858	35, 052	46. 187	15.872	16.376	25.572	24. 460	29.698
50	NT2RP2003034	97.685				40.853	44.346	20, 833	60.360
	NT2RP2003042	32.097	30, 146			14, 406	14.312	25, 483	23.898
								29.611	20. 940
	NT2RP2003050	43.965	23.480			15.913	20. 938		
	NT2RP2003060	43, 467	23, 385	32.696	13.554	17.473	48.442	37.686	31.235
	NT2RP2003073	90.622					68.737	36. 287	64.071
	CHIERLEGOSOLS	30.022	1. 7. 930	1 000.010	1 -334			<u> </u>	

497

Table 88

		** ***		105 001	00 000	00 100 1	CA SEA T	20 115 1	45 353
	NT2RP2003099	69.980	61.964	197.831	28.962	29. 485	52.756	36.145	46.753
	NT2RP2003108	22.037	23. 450	29,734	12.784	12, 243	25. 414	19.582	14, 441
			76.490	219.003	26.090	53, 025	89. 403	96.086	53.165
5	NT2RP2003115	175. 202							
3	NT2RP2003117	132. 572	135. 105	428.449	65. 631	66.802	77.649	41.504	75.169
	NT2RP2003121	77. 521	49.860	42.009	15.143	26.745	31.652	32.041	27. 916
		35. 377	29.656	27.135	9.957	16, 383	12.805	20. 265	8, 252
	NT2RP2003125								
	NT2RP2003127	29. 566	16.867	20. 397	5. 212	10.531	18.240	19.752	7. 540
	NT2RP2003129	50. 461	54. 112	157.477	25.025	29.892	16.686	23. 103	33.770
	NT2RP2003137	8.001	18.759	14, 140	10.321	7.469	15, 281	5. 429	3. 225
10							30.612	24.709	34. 031
	NT2RP2003138	52. 296	44. 278	85.267	21.446	22. 168			
	NT2RP2003146	55. 329	37.398	52.403	14. 492	12.222	29.608	23. 329	32.663
	NT2RP2003148	150. 386	104. 523	330.270	60.524	70.523	90.836	76.602	100. 291
	NT2RP2003150	26, 432	11. 157	23.761	15.678	11, 132	36.468	7.133	18, 954
						23, 422	50.314	42.129	
	NT2RP2003157	58. 172	46.518	64, 963	42.288				48. 145
	NT2RP2003158	44. 248	20. 906	37.740	8.136	17.954	27.119	19.062	38. 471
15	NT2RP2003161	19.274	11.968	16,062	2.701	7, 578	17.086	7.441	31.024
		49, 401	19.110	28.830	12.219	12,819	22. 155	19.787	34, 090
	NT2RP2003164								
	NT2RP2003165	89. 985	65. 955	218.487	37.132	35. 205	34. 406	24.887	33, 303
	NT2RP2003177	43. 596	22.142,	51.196	11.148	3.934	15.303	13.349	69.154
	NT2RP2003179	69, 718	46.328	169.618	30.883	22.456	37.444	43.967	45. 776
				22.293	13.420	10.852	20. 144	19.065	43, 611
	NT2RP2003194	144. 137	17. 980						
20	NT2RP2003206	7.840	5. 389	10.850	5.014	4.029	11.290	7.725	3.709
	NT2RP2003210	51. 322	21.586	38.521	12.974	17.884	37.608	30.477	29.805
	NT2RP2003227	42. 906	18.716	24, 162	17, 143	9, 513	37.425	15. 949	23, 165
		58.612	29. 572	62.903	22.926	28.577	30.449	37.367	63, 378
	NT2RP2003228								
	NT2RP2003230	5. 885	10.431	148.181	5. 253	9. 252	9.617	6. 228	22. 492
	NT2RP2003231	69. 197	41.691	59.459	34.789	15. 272	58.827	33.617	37.859
0.5	NT2RP2003237	30.563	38.860	123.572	28.832	11.050	15. 189	9. 580	23.097
25		33.469	21.053	50.845	20. 348	11.513	25.692	7.484	35, 924
	NT2RP2003239								
	NT2RP2003243	145. 467	34. 182	76.360	17.705	28.702	66.482	55.093	28. 921
	NT2RP2003265	29. 516	23.976	32.673	9.710	15.918	17.608	20, 157	14.165
	NT2RP2001267	65.087	29.515	67.969	24. 282	21.518	34, 797	27. 241	43.679
	NT2RP2003272	41. 457	22, 351	19,055	27.076	19.762	28.028	26. 982	45, 977
								26. 460	
30	NT2RP2003277	107. 913	82.634	92.986	31.633	32.424	67.812		53.116
00	NT2RP2003280	19, 151	14.918	20.689	11.633	7.567	43. 338	5. 070	12.961
	NT2RP2003286	21.848	17.740	29.829	11.104	6.965	28.110	26.734	25. 233
	NT2RP2003293	94.719	83.407	364.260	76.134	56.105	78. 539	44. 376	97.047
							15.088	25.617	15. 166
	NT2RP2003295	17.874	16.886	18.717	18. 256	19.625			
	MT2RP2003297	9. 592	10.816	15, 547	2.211	5.615	8.461	10.162	5.662
	NT2RP2003300	15, 144	16.953	26.519	1 10. 354	14.045	6.847	8.974	11.058
35	NT2RP2003302	22.071	15.550	64, 230	26.397	10.289	12.880	11.722	68. 523
	NT2RP2003307	22. 086	9.418	17.120	5. 220	6.112	15.691	17. 396	7.096
	NT2RP2003308	17. 436	24.315	20.930	11.886	7.814	20. 422	12.860	31.766
	NT2RP2003311	22.001	9. 144	13.842	5. 360	10.074	18.616	5. 176	21. 146
	NT2RP2003329	44.872	14.471	19.961	10. 976	13.401	22. 292	12.093	14.770
	NT2RP2003339	20.422	19. 625	85.412	16.458	12.443	17.818	9. 125	13.152
40	NT2RP2003345	23. 118	8. 297	17. 237	4.695	8.379	12.952	12. 259	23.215
40									
	NT2RP2003347	12. 189	4.636	9.822	7.720	7.500	12.461	7. 182	16.011
	NT2RP2003367	10.794	19.358	21.160	7.884	14.120	12. 142	14.419	13.409
	NT2RP2003369	41.141	18.327	38.318	11.072	14, 355	33, 971	28. 126	19.613
		55, 891	32.218	76.058	21.558	27.536	76.861	50.564	36. 175
	NT2RP2003383								
	NT2RP2003390	73.620	57.765	91.034	41.124	35.539	63.744	46.234	42.766
45	MT2RP2003391	241.564	161.239	277.051	75. 828	95. 432	220.668	152.546	143.981
	NT2RP2003393	11.758	13.507	20.112	4. 587	11.809	12.940	19.991	21.749
	NT2RP2003394	7. 323	9.816	9, 506	2.871	10.713	1. 307	6.346	14.753
	NT2RP2003401	25. 259	3.938	8.376	2.832	4, 096	7.246	16. 169	7.442
	NT2RP2003403	31.239	26.205	109.072	18.680	14. 206	9. 380	14. 946	8.745
	NT2RP2003433	79.603	33.408	70.460	19, 431	29. 526	42.730	34.783	28.629
	NT2RP2003445	38. 525	33.248	95,090	23.648	21.333	27.951	21.347	33.662
50									
	NT2RP2003446	67. 228	39.971	49.302	18.878	21.829	54. 339	39. 113	29. 464
	NT2RP2003456	1. 902	13.833	10.178	7.437	1.522	5.049	1.410	3. 486
	MT2RP2003466	72.001	27.022	47.862	12.506	26.814	66.543	51.004	41.515
	NT2RP2003469		29. 791	90.766	19.568	17.254	24.857	16. 952	39.575
		35. 915							
	NT2RP2003470	20.820	31.916	84.744	64.680	20. 126	61.522	22. 215	98.657

498

Table 89

55

								- 3 300 1	10 010
	NT2RP2003471	7. 424	5. 547	6.488	7.037	5. 447	6. 505	7.782	10.212
		78.094	65.408	137.798	31.787	40.594	58.633	37.776	39.678
	NT2RP2003480				7.870	5.725	11.076	8.329	14, 404
-	NT2RP2003495	15. 982	11.924	14. 233					
5	NT2RP2003499	55. 449	<b>13.382</b>	25. 597	4. 229	14. 517	54. 430	36.252	15, 105
		55. 425	27.024	46.996	11.964	7.933	31.002	31.997	27. 989
	NT2RP2003505					12. 205	23, 185	12.152	24, 906
	NT2RP2003506	29.029	19.815	26.696	9.949				
	NT2RP2003511	85. 237	37.479	50.383	22. 212	25. 152	50.854	41.079	36.551
			4. 521	4, 122	3.531	5.027	3,740	2.918	7.377
	NT2RP2003513	2.085					30.516	43.651	39, 873
	NT2RP2003517	37.834	17.587	35, 502	11, 597	12.069			
10	NT2RP2003522	24. 832	37.794	30.938	13.985	21.613	21.384	15.975	15.713
			77.947	318,616	53, 968	64, 300	64.511	45, 220	44.281
	NT2RP2003525	112.839				35.543	46.891	33, 401	37,402
	NT2RP2003533	95. 494	87. 932	267.080	44.833				56.566
	NT2RP2003541	59. 237	40. 256	51.598	18.653	24. 451	41.018	38.504	
		60.456	24, 016	25.882	11.661	16.145	17. 623	31.288	25.312
	NT2RP2003543			11.338	12, 197	5.950	2.774	8.060	34,030
	NT2RP2003545	5. 111	9. 859				24.477	17. 350	31,685
15	NT2RP2003559	26. 905	22. 287	37.874	13.292	12.911			
	NY2RP2003564	29, 146	18.045	64.896	13.749	13.213	15. 703	17.055	25.744
			106.907	131.344	34.826	44.614	78.728	62.826	61,650
	NT2RP2003565	71.340					61. 162	50. 325	46.459
	NT2RP2003567	70.892	54. 381	72.715	19.440	21.958			
	NT2RP2003575	8.045	11.848	16.656	3.697	4. 227	5. 271	7.753	9,628
		94, 175	119, 128	189.789	159.528	39, 210	94. 530	84. 153	280.017.
	NT2RP2003576					32.853	121. 326	99.589	58.803
<b>20</b>	NT2RP2003579	55. 985	110, 923	72.170	19.865				
	NT2RP2003581	72.231	34. 935	63.218	15.922	25. 161	44. 829	45.801	38.825
	NT 2RP2003587	109.102	46, 403	76.235	20.483	28.667	127.344	62. 139	47.892
				26. 653	9.837	5.016	24.313	17.397	36, 147
	NT2RP2003590_	27.361	26. 330						
	NT2RP2003593	98.848	66, 189	91.401	17.565	31.030	61.583	54. 982	56.233
	NT2RP2003596	20. 156	17.830	46. 567	15, 376	7.364	8, 849	10.462	35.925
				53.708	30.551	33, 831	64. 394	76.259	72.122
25	NT2RP2003599	99. 163	72.506					30.947	25.783
	NT2RP2003600	39.566	25.200	27.397	13.373	16.019	22.567		
	NT2RP2003604	30, 188	48.497	24.769	15.941	13.513	20.832	18.908	35.739
			10.012	13.520	5.134	7.235	8.896	12,558	21, 197
	NT2RP2003629	12.593				24.270	37, 506	32.166	28.383
	NT2RP2003530	55.769	31.553	55. 456	13.290				
	NT2RP2003643	20. 532	14.638	38.212	9.363	17.760	18.713	18.506	19.629
	NT 2RP2003655	46.795	29,612	38. 397	10.145	18.688	20. 220	24. 997	18.685
30				21.831	11.981	11.047	39.022	14.701	15,715
	NT2RP2003664	23.372	28.188					46.022	49.968
	NT2RP2003668	98.074	77.678	215.011	48.838	58.733	45. 358		
	MT2RP2003687	36.469	27.937	30, 101	11,600	12.659	14.676	15. 349	16, 155
				140. 266	28.579	24.877	10.915	18.651	30.704
	NT2RP2003691	57.166	65.814						35, 262
	NT2RP2003702	77. 231	74.259	157.835	37.740	29.269	33. 935	36.174	
	NT2RP2003704	33.958	19.273	90, 406	13.087	15.614	12. 526	13.208	27.631
35			9.802	10, 782	1.905	1.888	20.850	8.045	6.106
	NT2RP2003706	15.581					11. 248	12.533	12.834
	NT2RP2003713	16.960	13.155	19.058	12. 333	6.597			26.140
	NT2RP2003714	58, 106	48,190	156. 974	28.216	25, 935	21.990	15.804	
	NT2RP2003727	16.878	30.048	11.471	24.840	10, 360	26. 581	2.051	18.209
			27.626	24.696	15.279	8.490	48.230	26.577	18.778
	NT2RP2003737	35.097					15.115	11.036	11.385
	NT2RP2003751	24. 927	12.926	14. 285	5.654	5. 362			
40	NT2RP2003760	61.984	14.851	34, 689	31.937	11.912	70.013	35. 412	50.086
	NT 2RP2003764	70.923	28.030	49.140		33.253	31.845	28.042	21.978
						10.396	11.852	16, 178	10.912
	NT2RP2003769	42.617	20.886	27.599				43.465	55.063
	NT2RP2003770	137.506	66.296	82.283		19.657	59. 586		
	NT2RP2003777	79.392	37.432	49, 453	21.542	23.944	31.481	38.443	30.003
						41.064	65.158	51.558	43.936
	NT2RP2003781	113.598	78.822					21.078	32.965
45	NT2RP2003785	39.008	38.895			81.398	60.210		
	NT 2RP2003793	29.403			11.279	11.070	27.094	13.519	16, 114
						57.427	54.142	52.055	74.576
	NT2RP2003806	141.377							115, 589
	NT2RP2003825	200.861	142.661			83, 143			
	NT2RP2003840	100.905			27.801	38.812	73.708		43.672
							109. 107	66.696	63.138
	NT 2RP2003857	135. 915							
50	NT2RP2003859	112, 898							
	NT2RP2003871	16, 891	14.873	18.946	20.075				
		20.553					22.067	11.629	10.917
	NT2RP2003876								
	NT2RP2003878	10.935							
	NT2RP2003885	86.861	91.093	40.63	9.621				
	NY2RP2003898	42.684			13.576	37. 187	19.007	22.509	33. 529
	[M16KFEDU3838	72.004	1 30. 30						

Table 90

			104 005 1	100 475	46 004 T	48.594	124. 353	51.962	58. 344
	NY2RP2003902	147. 643	124. 985	109.475	45. 984	109. 998	129. 880	47.537	95. 222
	NT2RP2003912		242.124		29. 243				3, 142
-	NT2RP2003931	26.887	8.179	6.459	2.307	5. 260	8. 153	1.858	
5	NT2RP2003940	185. 397	64.618	262.034	55, 607	30.649	41.635	23. 343	65.087
	NT2RP2003950	36. 158	19, 195	49.413	13.592	20.939	19.343	26.770	21.989
	NT2RP2003952	15. 955	17, 931	35,750	13.974	12.405	27.300	20.083	13.016
	NT2RP2003968	45. 877	22.833	13.459	11.361	12.355	12.353	12.010	25, 113
		37. 958	44, 808	95.495	38.986	28, 544	21.209	8.325	15. 117
	NT2RP2003976		43, 006	57.657	15. 338	29. 345	30. 659	23. 563	25.867
10	NT2RP2003981	38.654			16. 394	44. 914	84. 097	45. 289	33. 280
	NT2RP2003984	132.353	65.644	60.516			70.656	43.927	53, 945
	NT2RP2003986	186.062	146.313	421.324	109.891	71.468	64. 387	44, 174	58. 384
	NT2RP2003988	112. 131	82.329	348. 163	81.784	60. 909			38, 113
	NT2RP2004013	35. 821	31.054	41, 104	24.447	20. 809	33.899	21, 394	
	NT2RP2004014	51.068	77.076	125. 407	38.647	29. 948	34. 055	26.943	33.783
	NT2RP2004036	34. 592	12.491	12.862	9. 166	7.965	9, 771	12.722	18. 319
15	NT2RP2004041	61.828	31,728	66.443	16.578	28.668	39.049	31, 113	30. 197
	NT2RP2004042	95. 416	34, 528	56.458	18. 193	31.581	50. 180	28.757	19.510
	NT2RP2004049	30.836	31, 163	33.858	10.780	19. 423	28. 518	29.763	8. 339
	HT28P2004060	33. 939	22.080	47.086	13, 117	10.598	29.819	24.922	24.074
		36. 939	51.977	61.500	23. 281	20.470	26,729	15, 403	25. 483
	NT2RP2004066		33.889	47.332	22.168	14.676	23.715	30.550	18, 563
20	NT2RP2004059	29. 217		35. 232	4, 198	9.970	5.069	6.316	20.634
20	NT2RP2004076	9. 020	12.153		4. 309	8. 489	27.512	5. 327	10.188
	NT2RP2004080	23.022	8.835	21.995		33.602	10. 431	18. 338	56.090
	NT2RP2004081	38. 786	30.091	83.806	31.063		26. 497	22.648	13.621
	NT2RP2004098	47.764	21.424	36.354	14.003	22. 548		20.636	48, 103
	NT2RP2004108	28.744	38.559	67.714	34.947	23.442	39.884		
	NT2RP2004124	43.031	24.659	37.232	12.008	12. 194	23. 487	10, 186	21.361
25	NT2RP2004130	62.738	36.522	73.772	37.407	24. 390	44. 094	20.478	34. 479
	NT2RP2004133	163.939	56.278	112.008	40.808	61.092	157. 167	95. 384	52.343
	NT2RP2004141	49. 570	22.611	50.916	9.793	20.924	53. 203	22.033	30. 466
	NT2RP2004142	34. 850	23.492	33.078	17.102	15. 132	27.703	11. 237	17.601
	NT2RP2004152	14. 256	11.207	21, 943	19.655	8.860	14.997	12.981	8. 353
	NT2RP2004185	147.447	92.813	238.228	40.497	54. 357	70.413	30.081	44. 940
	NT2RP2004170	107.111	64.978	194.673	41.028	56.020	66. 291	58.470	56, 553
30	NT2RP2004172	22.440	15.213	19.562	6.795	12.099	15. 400	14.334	12.024
		120. 902	23.723	54.734	12.552	24.966	70.512	39.664	28. 280
	NT2RP2004176		30. 327	45.178	12.821	11.733	33. 905	35. 842	30, 011
	NT2RP2004179	72. 406	21.870	33.704	11.364	19.908	8. 982	12.208	16. 442
	NT2RP2004187	25. 235		37.882	8. 251	10.063	16.897	16.826	36.649
	NT2RP2004190	33, 406	32.037		35. 398	25. 386	70.700	59. 372	84.014
35	NT2RP2004194	84.064	81.541	54.017		35. 795	83.939	40. 164	46.168
	NT2RP2004196	105.711	65, 320	61.236	35. 178		102.336	55. 538	55. 936
	NT2RP2004205	144.445	71.761	300.198	38.897	46. 886		17. 403	14. 550
	NT2RP2004207	34.894	12.571	14.703	6.333	7.074	34.908	27. 469	21.672
	NT2RP2004226	63.802	26.160	69, 559	17.665	24. 160	72.242		
	NT2RP2004232	19.053	14.404	25.695	7.555	9.877	15. 593	12.523	32.679
	NT2RP2004239	49.739	30.594	47.640	22.915	18.596	31.416	32.672	84. 520
40	NT2RP2004240	43.946	56.977	36.742	39,656	38.450	39.881	22.758	41.302
	NT2RP2004242	24, 272	10.675	24. 496	11.743	14.023	31.038	18.900	15. 124
	NT2RP2004245	18.673	23.813	15. 945	12.936	16.016	18.326	7.178	10.903
	NT2RP2004270	234. 182	227.894	511.563	104.046	110.474	124. 225	90. 436	89. 248
	NT2RP2004300	59.573	43.407		15.466	13.124	34.892	25.094	19.570
	NT2RP2004304	30. 539			13.187	14.829	18.430	12.663	17.214
45	NY2RP2004313	52,639				13. 307	42.833	29.621	25.693
		7.937					5. 139	1.817	5.009
	NT2RP2004316	16.873						12.417	8.754
	NT2RP2004321							18.808	17.712
	NT2RP2004336	27.640						70.845	110.855
	NT2RP2004339	253.896							12.471
	NT2RP2004347	39.311							23.600
50	NT2RP2004364	71.148							
	NT2RP2004365	27. 548							27.122
	NT2RP2004366	34. 341							
	NT2RP2004373	28.456							14.963
	NT2RP2004375	22.258							28.153
	NT2RP2004389			3 17.94	11.246	10.83	22.718	14.078	16.693

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Table 91

		00 000 1	136 436 [	185. 407	07.306	71, 728	98.742	40, 421	94. 207
	NT2RP2004392		136.238	232.453	39. 577	40. 329	51.827	19.795	36. 180
	NT2RP2004396	74. 685	\$5.569	62.661	13. 504	14.626	15.041	17.402	11. 134
5	NT2RP2004399	60.880	42. 455		31, 390	26. 256	16.692	21.998	27.979
3	NT2RP2004400	48. 188	46. 127	127.225	33, 584	39. 340	32. 995	41.822	41. 552
	NT2RP2004404	94, 197	59.189	80.085		73, 546	51.855	24.894	53. 454
	NT2RP2004410	42. 321	76.331	55. 926	19.723		29.062	12.037	4.016
	NT2RP2004412	13.609	18.755	18. 039	11.352	6. 207	35. 340	12.897	8. 527
	NT2RP2004414	14.965	13.344	29.690	8.080	8. 676	4. 256	11.714	5. 665
	NY2RP2004425	15.759	4.692	13. 145	5. 794	4. 150	15.766	15.563	25. 352
10	NT2RP2004447	42.510	30.709	103. 682	26. 465	17. 475	65.475	55.192	37. 759
	NT2RP2004463	64.696	47. 400	81.626	29. 385	29. 125	26.270	24. 224	25. 991
	NT2RP2004476	27.281	77.743	30. 875	42.538	9.672		12.795	25. 305
	NT2RP2004488	22.602	16. 334	32.445	12.940	12.612	19.801	48.408	47.844
	NT2RP2004490	108.056	33. 325	36.585	11.778	28. 508	83.898 24.630	15.828	7.643
	NT2RP2004495	24. 445	8. 305	18.686	11. 202	4. 044			14. 259
15	HT2RP2004512	4. 285	7, 813	16.614	6. 915	11.355	6.603	2.640	44. 602
	MT2RP2004523	100. 195	69. 639	192.670	43, 236	39, 566	47. 481		
	NT2RP2004524	44. 944	32, 536	60.310	17. 428	15, 331	26. 455	22.167	50. 697
	NT2RP2004536	61.814	19. 213,	31, 957	8.029	18. 302	52.061	24.818	16.740
	NT2RP2004538	844. 732	696.798		422. 320	403.488	580. 281	434. 455	470.608 58.770
	NT2RP2004548	81.639	84.667	179.445	54. 320	34.612	101.391	35. 028	4. 857
20	NT2RP2004551	20, 101	20. 257	8.701	5. 567	6. 509	4. 732	2,996	107. 235
	NT2RP2004556	186.686	124. 741	397. 345	91.884	102.226	91.039	70.486 45.819	146.073
	NT2RP2004568	92.661	117.910	131.215	47.958	44.000	46. 192		55. 832
	NT2RP2004580	117.798	112.312	308.956	61.075	41.911	54. 139 75. 503	28.004 38.854	53. 952
	NT 2RP2004585	88. 489	51.782	72.459	31.850	12. 237		2. 284	3. 479
	NT2RP2004587	9.681	12.544	13.758	5. 129	6. 286	5. 708 3. 168	5. 020	19. 533
25	NY2RP2004594	17.013	7. 543	15, 550	11.674	7. 962	5.752	8. 192	20. 142
	NT2RP2004600	24. 043	10. 196	26.881	6.520	4, 919	36.752	5. 232	36. 380
	NT2RP2004602	123.606	61.805	80.505	32. 526	37. 163	58.545	65.119	58.082
	NT2RP2004606	95. 195	78.770	115.775	31.102	36, 965 25, 720	49.230	34.724	39. 520
	NT2RP2004614_	88.734	53. 501	57. 570	36.772	8. 537	50. 158	15, 799	9, 179
	NT2RP2004648	20, 700	23.018	14.031	7.353	6,707	24. 083	10.703	5.977
30	NY2RP2004655	15. 547	12.030	20. 925 45. 941	18. 159	13, 692	93.784	43.213	29.634
••	NT2RP2004664	115.653	30. 969		8.337	17.064	23. 260	22.585	18.670
	NT2RP2004670	37. 342	20. 435	29.733	52. 918	33, 597	43. 245	31.102	40.195
	NT2RP2004675	90. 376	87.838	71.220	24. 851	34, 241	54. 143	45.414	29.175
	NT2RP2004681	80.974	6.449	9.318	5. 269	6. 188	5. 655	17.368	7, 173
	NT2RP2004689	15. 361	57.745	95.083	23. 386	38. 263	34,748	18, 462	31.462
35	NT2RP2004709 NT2RP2004710	76.835 55.266	57.910	39. 262	18.404	10.078	36.682	30.725	36, 367
	NT2RP2004710	326. 635	50.412	98.334	21.234	65.676	230.530	162.452	35.853
	NY2RP2004736	151, 717	95. 950	265. 487	84.538	82.942	67.704	64.264	123.565
	NY2RP2004743	34, 118	25, 149	128.802	17, 805	15.041	28.540	44.641	29.720
	NT2RP2004750	83. 958	75. 396	199.356	68.993	52.468	133.541	50.743	56.041
	NT2RP2004755	31.604	24. 450	46.432	13.888	69.303	26.643	15.757	22.713
40	NT2RP2004757	79.661	59. 962	217.503	30.858	29. 576	29.740	25. 153	35. 482
	NY2RP2004768	13. 287	13.098	19.823	9.173	5. 193	3, 545	2.323	8.664
	NT2RP2004775	10. 197	8.827	40.973	5.720	4,909	3.010	5.098	1.954
	MT2RP2004791	68, 964	37, 186	133.612	23.163	25. 209	12.978	21.406	22,080
	NT2RP2004794	230. 935	115.789	236.516	45. 963	115, 577	229. 430	167.093	66.975
	NT2RP2004795	38.086	12.315		9.762	10.237	23.540	30.190	27.839
45	NT2RP2004799	32.524	12. 267	12.671	2. 945	22.824	24. 117	5. 268	5.775
.5	NT2RP2004802	10.030	10.579	12.121	10.897	8.541	5.714	8.012	10.032
	NT2RP2004810	42.256	25, 180	28. 300			15. 976	14, 419	10.508
	NT2RP2004816	30. 283			17.849		23.062		21.647
	NT2RP2004837	247.337					220. 470		
	NT2RP2004841	18.863						6.007	
50	NT2RP2004847	273. 546	127.737				209. 860		
30	NT2RP2004861	39.358	31.567				19.568		
	NT2RP2004897	15. 367					26. 917		
	NT2RP2004932	183.953							
	NT2RP2004933	18.560							
	NT2RP2004936	10.618	16. 16:	27.370	5. 543	8. 959	13. 920	6.220	8.621

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Table 92

								34 477
NT2RP2004951	30, 413	16.712	16.279	18.835	12.085	15.888	11.101	14.477
NT2RP2004959	7.613	10.358	13, 406	5.314	5. 926	11.986	2.543	5.752
	42.335	32. 379	69.235	34. 253	21.447	34.663	18.456	42. 255
		30.353	89. 154	14.113	9. 384	17.622	8.128	20.787
NT2RP2004962	30.669						22.899	25.847
NT2RP2004966	42.472		27.864	9, 661	13.817	26.018		
NT2RP2004967	57. 426	40.541	179. 390	31.892	23.923	31.052	16.791	43.678
NT2RP2004974	31.596	11.054	27, 118	11.874	12.196	35. 458	18.873	24, 149
		58. 297	58.744	11.187	26.598	42.390	34.073	15, 958
NT2RP2004978	92.366			2. 288	3,775	3. 554	3.062	0.000
NT2RP2004982	2.062	5, 171	6.063					62.719
NT2RP2004985	87.939	67.149	78.678	45.629	27.293	69.956	48. 241	
NT2RP2004999	54, 349	44. 327	160, 162	26.886	23.352	26.240	26.943	41.559
NT2RP2005000	26.080	14. 589	21, 728	7.864	8,002	19.702	12.179	15. 480
		13. 183	23,055	6, 161	9.633	14.650	18.615	13. 44
NT2RP2005001	26.862			39.371	25. 182	33, 952	24.278	47.01
NT2RP2005003	69.867	63.795	165.289				24. 522	29. 43
NT2RP2005012	30.982	21, 105	42.355	15.018	14.157	41.891		
	111.833	49. 415	78.251	22. 107	42.271	59. 226	38.060	18.69
NT2RP2005020	60.906	32.923	38. 225	11.918	20.379	16.776	20. 985	35.43
			37. 383	8.777	13.169	17.643	25.803	22.97
NT 2RP2005022	44.931	25, 614			17, 117	35. 304	31.116	36.53
NT2RP2005027	57.511	85.851,	98. 132	22, 401				3.24
NT2RP2005031	14,601	8, 758	14.468	5. 468	5, 699	7.564	8.732	
NT2RP2005035	61.937	41.750	49.801	22.387	27. 920	58. 127	29. 585	39.14
WYANDONAFOOT	27.745	16. 434	26. 221	9. 584	20.837	24.795	26.368	30.42
NT2RP2005037			12.702	3.787	5. 660	9.747	35. 202	6.79
NT2RP2005038	13.976	3. 551				30. 149	28.713	25.89
NT2RP2005048	55.851	47. 103	55.038	22.550	27.846			
NT2RP2005069	89. 545	160.853	309.743	119.361	135.285	158.356	127.275	142.12
NT2RP2005073	28.642	24.071	29.062	8.191	14.897	17.052	25.028	53.37
		11.744	11, 103	4.196	6.885	7.430	12.482	11.24
NT2RP2005097	17.446			4.736	8, 210	16.355	10.080	48.38
NT2RP2005108	22.062	6.419	8.005				122.651	68.89
NT2RP2005116	161.700	67. 851	96.374	39.093	51,697	82.025		
NT2RP2005126	24,712	30.925	25.757	24. 268	22.706	35. 722	14.976	36. 43
NT2RP2005135	38.054	16.075	22.834	7. 220	8.729	25.814	9.825	8.45
	25. 339	21.341	24.789	9. 299	9, 331	10.389	15, 907	13.63
NT2RP2005139			18.762	7.827	14.629	21.623	15.226	7.6
NT2RP2005140	25. 302	14. 152					25.702	14.4
NT2RP2005144	57.910	24.627	35. 294	9, 403	20.129	22.753		
MT2RP2005147	35. 344	15.053	40.777	7.320	13.980	7.943	9.818	7.0
NT2RP2005148	71.460	50.351	93, 151	24.862	24. 403	40. 037	28. 927	30.9
NT2RP2005159	32.863	9, 249	11.688	11.160	6. 240	11.164	8.584	7.6
	33.677	20.731	31.783	9.893	9. 733	6.520	16.473	12.8
NT2RP2005162				125. 386	135. 331	256.832	253.752	198.4
NT2RP2005163	406.419	245. 982	312.290				20.783	14.2
NT2RP2005168	44. 795	9. 276	16.080	8.798	9, 082	15.704		
NT2RP2005181	58, 670	24.911	19.589	19, 590	10.885	16.528	28.301	18.9
NT2RP2005204	51.862	36.997	48. 257	21,014	21.820	26.400	20.001	33.9
		44.601	71.232	24.297	39.166	94, 145	83.743	57.0
NT2RP2005219	118. 951			37. 680	26. 287	29.966	35, 172	44.3
MT2RP2005227	63. 965	85.586	198.792			91.954	81.365	194. 9
NT2RP2005237	95. 186	85.568	117,090	33.460	57.400			
NT2RP2005239	45, 116	20.823	33. 169	16.031	8.498	11.991	27.107	20.6
HT2RP2005247	55, 177	33.524	91,868	28. 505	25.628	27.978	38.659	30.3
NT2RP2005254	67.776	32,943	35, 931	20, 251	16.723	35.298	24. 338	25.
	36.792	20.989	23,940	12.941	20.407	34.731	24.269	12.4
NT2RP2005270				15. 429	19.430	30.784	9.484	44.1
	34.791	50.008	19.917			23.753	21.837	56.
NT2RP2005276	75. 555	49, 491	25. 557	16. 229	7.738			
NT2RP2005287			1 16 471	9.015	8,671	29.745	15.169	27.0
NT2RP2005287	84, 486	30.764	16.471					723.
NT2RP2005287 NT2RP2005288	84, 486		195.975	30. 937	28.926	33. 261	20.112	
NT2RP2005287 NT2RP2005288 NT2RP2005289	84, 486 74, 343	79.634	195.975	30. 937	28.926	33. 261 25. 911	15.734	9.
MT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293	84, 486 74, 343 32, 574	79.634 17.527	195. 975 13. 426	30. 937 14. 326	28. 926 12. 226	25. 911		9.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315	84, 486 74, 343 32, 574 30, 488	79.634 17.527 58.065	195. 975 13. 426 52. 471	30. 937 14. 326 14. 353	28.926 12.226 24.658	25. 911 23. 599	15.734 29.610	9. 56.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005322	84, 486 74, 343 32, 574 30, 488 54, 278	79.634 17.527 58.065 66.487	195. 975 13. 426 52. 471 69. 926	30. 937 14. 326 14. 353 39. 640	28. 926 12. 226 24. 658 44. 675	25. 911 23. 599 56. 756	15.734 29.610 45.067	9. 56. 30.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005322	84, 486 74, 343 32, 574 30, 488	79.634 17.527 58.065	195. 975 13. 426 52. 471 69. 926 114. 652	30. 937 14. 326 14. 353 39. 640 20. 676	28. 926 12. 226 24. 658 44. 675 66. 949	25. 911 23. 599 56. 756 180. 520	15.734 29.610 45.067 130.512	9. 56. 30. 46.
MT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP20053315 NT2RP2005322 NT2RP2005325	84. 486 74. 343 32. 574 30. 488 54. 278 244. 369	79.634 17.527 58.065 66.487	195. 975 13. 426 52. 471 69. 926 114. 652	30. 937 14. 326 14. 353 39. 640 20. 676	28. 926 12. 226 24. 658 44. 675	25. 911 23. 599 56. 756	15.734 29.610 45.067 130.512 34.194	9. 56. 30. 46. 55.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005322 NT2RP2005325 NT2RP2005336	84, 486 74, 343 32, 574 30, 488 54, 278 244, 369 118, 767	79.634 17.527 58.065 66.487 45.065 75.218	195. 975 13. 426 52. 471 69. 926 114. 652 151. 013	30. 937 14. 326 14. 353 39. 640 20. 676 52. 856	28. 926 12. 226 24. 658 44. 675 66. 949 19. 178	25. 911 23. 599 56. 756 180. 520	15.734 29.610 45.067 130.512	9. 56. 30. 46. 55.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005312 NT2RP2005322 NT2RP2005325 NT2RP2005326 NT2RP2005336	84, 486 74, 343 32, 574 30, 488 54, 278 244, 369 118, 767 83, 426	79.634 17.527 58.065 66.487 45.065 75.218 73.474	195. 975 13. 426 52. 471 69. 926 114. 652 151. 013 185. 631	30. 937 14. 326 14. 353 39. 640 20. 676 52. 856 30. 816	28. 926 12. 226 24. 658 44. 675 66. 949 19. 178 16. 652	25. 911 23. 599 56. 756 180. 520 13. 614 38. 395	15.734 29.610 45.067 130.512 34.194 18.655	9. 56. 30. 46. 55.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005322 NT2RP2005325 NT2RP2005336 NT2RP2005344	84, 486 74, 343 32, 574 30, 488 54, 278 244, 369 118, 767 83, 426 13, 456	79.634 17.527 58.065 66.487 45.065 75.218 73.474 15.006	195. 975 13. 426 52. 471 69. 926 114. 652 151. 013 185. 631 16. 224	30. 937 14. 326 14. 353 39. 640 20. 676 52. 856 30. 816 6. 558	28. 926 12. 226 24. 658 44. 675 66. 949 19. 178 16. 652 7. 385	25. 911 23. 599 56. 756 180. 520 13. 614 38. 395 8. 066	15.734 29.610 45.067 130.512 34.194 18.655 8.800	9. 56. 30. 46. 55. 27.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005322 NT2RP2005325 NT2RP2005336 NT2RP2005344 NT2RP2005344 NT2RP2005344	84, 486 74, 343 32, 574 30, 488 54, 278 244, 369 118, 767 83, 426 13, 456 29, 998	79. 634 17. 527 58. 065 66. 487 45. 065 75. 218 73. 474 15. 006 29. 498	195. 975 13. 426 52. 471 69. 926 114. 652 151. 013 185. 631 16. 224 40. 925	30. 937 14. 326 14. 353 39. 640 20. 676 52. 856 30. 816 6. 558 17. 105	28. 926 12. 226 24. 658 44. 675 66. 949 19. 178 16. 652 7. 385 11. 916	25. 911 23. 599 56. 756 180. 520 13. 614 38. 395 8. 066	15.734 29.610 45.067 130.512 34.194 18.655 8.800 19.200	9. 56. 30. 46. 55. 27. 3.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005325 NT2RP2005325 NT2RP2005336 NT2RP2005343 NT2RP2005344 NT2RP20053547 NT2RP2005354	84, 486 74, 343 32, 574 30, 488 54, 278 244, 369 118, 767 83, 426 13, 456	79.634 17.527 58.065 66.487 45.065 75.218 73.474 15.006 29.498 179.788	195, 975 13, 426 52, 471 69, 926 114, 652 151, 013 185, 631 16, 224 40, 925 410, 980	30. 937 14. 326 14. 353 39. 640 20. 676 52. 856 30. 816 6. 558 17. 105 73. 329	28. 926 12. 226 24. 658 44. 675 66. 949 19. 178 16. 652 7. 385 11. 916 74. 840	25. 911 23. 599 56. 756 180. 520 13. 614 38. 395 8. 066 12. 479 81. 380	15.734 29.610 45.067 130.512 34.194 18.655 8.800 19.200 48.521	9. 56. 30. 46. 55. 27. 3. 17. 65.
NT2RP2005287 NT2RP2005288 NT2RP2005289 NT2RP2005293 NT2RP2005315 NT2RP2005322 NT2RP2005325 NT2RP2005336 NT2RP2005344 NT2RP2005344 NT2RP2005344	84, 486 74, 343 32, 574 30, 488 54, 278 244, 369 118, 767 83, 426 13, 456 29, 998	79.634 17.527 58.065 66.487 45.065 75.218 73.474 15.006 29.498 179.788	195, 975 13, 426 52, 471 69, 926 114, 652 151, 013 185, 631 16, 224 40, 925 410, 980	30. 937 14. 326 14. 353 39. 640 20. 676 52. 856 30. 816 6. 558 17. 105 73. 329	28. 926 12. 226 24. 658 44. 675 66. 949 19. 178 16. 652 7. 385 11. 916 74. 840	25. 911 23. 599 56. 756 180. 520 13. 614 38. 395 8. 066 12. 479 81. 380 31. 822	15.734 29.610 45.067 130.512 34.194 18.655 8.800 19.200 48.521	9.6 56.8 30. 46.6 55. 27.1 3. 17. 65.

Table 93 ·

	NT2RP2005378	275.722	60.663	120.794	35.912	75. 334	165. 512	90.853	98.444
			47.813	76.113	25. 253	31.400	92.500	35.776	47.335
	NT2RP2005391	150. 127			29.969	28. 518	49. 057	25. 746	34.105
_	NT2RP2005393	70.899	55. 424	140.116					11.408
5	NT2RP2005407	49. 576	20. 202	38.801	8.339	17. 993	20. 349	19. 728	
	NT2RP2005419	14.831	11.867	19. 565	9.795	8.679	10.513	8. 946	8.857
	NT2RP2005425	18. 167	59.599	35.636	25.050	15. 104	8. 153	9.614	51.727
	NT2RP2005429	59.197	19.497	39. 350	10.173	18.944	57. 213	14. 988	13. 492
	NT2RP2005436	79.164	77.083	60.113	36.736	34.134	54. 347	21. 339	40.541
	NT2RP2005441	13.042	15. 338	15. 762	8. 369	12.826	20.597	9.547	15, 936
10	NT2RP2005442	38.553	25.938	32. 259	17. 285	15. 576	32.634	33.798	38.091
	NT2RP2005444	71, 342	49.614	44. 203	32.335	34.594	66.817	40.260	65.040
		14. 907	15. 128	11. 162	5.959	22.081	9.421	11.234	15.739
	NT2RP2005453		70.504	365. 826	82.692	104.746	121.659	116.087	102.686
	NT2RP2005457	140.563		11. 247	8, 652	9. 030	17.490	6. 559	3.649
	NT2RP2005458	20. 125	11.007		20.532	31.448	29.345	25. 049	51.072
	NT2RP2005463	33. 251	29.837	73.818		13.341	13. 525	14. 209	18. 361
15	NT2RP2005464	15.800	16.043	35. 864	14.911		14. 681	6.572	9. 479
	NT2RP2005465	14.668	18.280	26. 584	6.257	10.356		42.443	7.644
	NT2RP2005472	16.851	25.760	9, 199	8.686	4. 966	40. 418		
	NT2RP2005476	46.416	52.525,	104. 203	20.584	20.782	24. 546	5. 316	32.360
	NT2RP2005490	61.983	24, 419	28. 345	12.864	15.040	12.501	22.637	19. 383
	NT2RP2005491	374.811	74.888	145, 408	24. 336	165.612	317. 177	231.269	69. 296
20	NT2RP2005495	31.802	17.805	29.680	11.830	10.557	8.912	14. 827	34. 592
	NT2RP2005496	148.755	112.441	375. 031	47.535	53.667	47.282	40. 191	44. 995
	NT2RP2005498	44.735	18.772	34. 164	9.402	20.468	26.500	17.998	17.049
	NT2RP2005501	40.853	37,008	48. 454	14.020	18. 699	36.333	14.886	19.992
	NT2RP2005506	90.354	86.898	75. 939	25.611	32. 147	174.626	79.478	131.787
	NT2RP2005509	49.249	30.854	40. 983	21.945	13.500	50.085	24.330	36.909
05	NT2RP2005514	27, 107	19.658	27.479	12.890	10.652	12.518	18.695	17. 325
25	MT2RP2005520	17, 919	21.654	27. 300	18.855	10.163	12.223	7.568	30. 261
	NT28P2005525	39.486	38.604	46.862	28.621	21.332	32.985	26.679	36.176
	NT2RP2005531	14,400	12.033	22.722	7.730	9. 380	14, 414	16.744	11.422
	NT2RP2005535	101.541	107.605	200.015	82.259	60.740	56.504	51.248	118.559
	NT2RP2005539	66.664	29.346	46.698	21.888	19.870	64.043	30.246	26.001
	NT2RP2005540	20.513	15.829	14.697	8.223	3.931	49.149	7.536	29.160
30	NT2RP2005541	64.709	41.297	53.989	27.868	23.974	31.435	25.336	31.933
	NT2RP2005549	32.008	17.222	22. 169	5.861	8.219	16.966	10.809	20.882
	NT2RP2005555	32.893	26.046	65.848	10.597	20.624	14.475	13.940	32.764
	NT2RP2005557	17.755	22. 321	31.949	8.994	15.581	5. 592	13.074	7.963
	NT2RP2005581	90.896	89.844	311.596	54.248	36.454	51.670	42.717	57.487
	NT2RP2005586	15.319	12.081	23.020	7.054	4.455	13.988	9.947	14.644
35	NT2RP2005597	70. 922	36,752	50. 127	12.506	18.474	43.281	28.038	27.738
	NT2RP2005800	57.039	36.730	42.297	19.089	22.952	20.349	26.429	35.687
	NT2RP2005605	89.117	41.403	109, 938	32.943	40.472	75.058	52.177	50.487
	NY2RP2005614	7.627	7.626	13,603	2.503	13.051	6. 276	5.809	8.317
	NY2RP2005620	42.734	21.553	33.023	9.850	14.899	31.978	27.521	25.649
	NT2RP2005622	17.770	22.460	29. 124	15.992	11.139	27.623	9.965	36.314
40	NT2RP2005632	14.999	31.771	43.031	12.307	17.618	13.899	11.335	15.678
	NT2RP2005635	49.456	30.521	47.412	10.091	23.056	33.511	25.653	30.736
	NT2RP2005637	12.810	11.271	23. 258	10.723	0.000	8.150	7.172	12.007
	NT2RP2005640	4.097	3.653	9. 894	0.840	1,980	8.957	6. 220	1.795
	NT2RP2005645	20.889	32.389	36.306	18.400	17,660	5.119	17.090	35.045
	HT2RP2005651	73.019	20.719	36.098	13.026	13.892	30. 207	10.624	40.618
4.5	MT2RP2005654	39, 235	27.889	43. 919		15.864	16.064	25.659	25. 595
45	NT2RP2005666	62.014	31.370	41, 680		18.813	69, 986	43.533	15. 230
	MT2RP2005669	64. 432	53.672	65. 910		25.429	55. 388	61.239	61.894
	NT 2RP2005670	37.363	15. 333	17. 547		14.756	35.642		14. 161
	NT2RP2005671	43.306	44. 120			17.143	63.049		21.799
	NT2RP2005675	142. 194	57. 967			42.418	100.132	100.664	78.669
			27. 395			10.519	19.049	11.915	16.611
50	NT 2RP2005683	25. 353					4. 338	11, 115	15.932
	NT 2RP2005690	15.846	16.544			24.945	11.950		28.108
	NT 2RP 20056 94	76.694	67. 508 185. 579			135.844	350.114	247.379	185.727
	NT2RP2005701	423.656	13. 221						16. 350
	NT2RP2005712	27.492							
	NT2RP2005719	10. 978	10.918	1 13.4/4	0.130	1 13.176	1 ,0. 700	1	7. 7. 7. 0

503

Table 94

	LUZABBOOK 233	24 666	61.425	70.544	51.843	34,010	57. 142	30.735	84.009
	NT2RP2005722	34.666						9. 230	34. 076
	NT2RP2005723	37.670	25.612	103.399	9.672	11.861	29.530		
	NT2RP2005726	84. 115	36.206	48.072	11.995	17. 484	39.045	38.061	22.448
5	NT2RP2005729	58.884	54.269	60.427	19. 257	22. 993	12. 151	26. 199	35. 691
	NT2RP2005731	17.800	7.315	9.355	4.076	7. 122	6.849	10.218	5.724
		135.853	80. 248	89.882	31.905	49. 498	82.876	94. 937	95. 379
	NT2RP2005732			192.481		56. 581	148.601	144. 906	98. 588
	NT2RP2005737	185. 624	120.622		48.397				
	NT2RP2005741	46. 137	31,647	35. 369	13. 164	19, 315	12.578	24. 931	17.774
	NT2RP2005748	37. 338	25. 300	30. 354	12.292	9. 999	24. 185	17. 843	16.711
10	NT2RP2005752	83. 285	59.855	77.223	35. 613	43. 031	19.000	35. 985	52.873
	NT2RP2005753	420.897	246. 480	444. 538	136. 522	121. 988	399. 581	356.877	181, 575
	DITOROGOGIZES	20.019	6.095	33.705	10.540	9, 232	5, 201	14. 128	11.843
	NT2RP2005763				10.684	15.614	27. 907	23. 447	10.054
	NT2RP2005767	46.813	15. 583	33. 205					
	NT2RP2005773	291.831	182.413	441.247	117. 268	110.788	192.144	163.936	144. 244
	NT2RP2005774	55. 239	48. 822	145. 962	59. 822	22. 432	33.644	24. 248	65. 283
15	NT2RP2005775	30.878	18.336	17. 192	11. 176	0.000	19.156	17. 205	15.094
	NT2RP2005781	56.648	31.034	24. 498	10.923	17, 115	16.751	30.579	26.075
	NT2RP2005784	153.655	51.631	100. 244	26. 189	25. 452	104.958	92. 590	20. 477
					24.721	19, 271	60.694	30. 122	14, 401
	NT2RP2005789	74. 249	51.916,	68.043					
	NT2RP2005799	71.863	10.045	12.797	6,316	3. 181	47.328	6.050	3.897
	NT2RP2005804	52. 496	43. 561	70.286	25.906	16.838	25.088	23. 482	31.711
20	NY2RP2005812	49. 420	17. 556	27. 165	8.036	15. 484	13.521	16.634	20, 990
20	NT2RP2005815	27.570	20.859	32.235	11.501	9. 452	14.728	19. 248	36.742
	NT2RP2005835	112.785	78. 188	150.766	35.828	53. 880	99. 576	62. 221	32.500
	NT2RP2005841	41.693	18.145	43.677	15.477	18. 203	8.667	17.036	33.652
				205. 381	30.242	23. 198	54.689	16. 871	24. 992
	NT2RP2005853	70.296	52.756					6. 163	12.771
	NT2RP2005857	23. 173	20.068	18.329	34.075	5, 778	4. 049		
25	NT2RP2005859	33. 168	17.202	37.200	12.544	13. 483	19.950	9.659	26, 739
23	NT2RP2005860	31.260	19.609	26.277	8.837	10.871	17.943	20. 399	13. 975
	NT2RP2005863	21.267	29.851	26. 528	17. 209	15. 572	12.614	18. 527	11.789
	NT2RP2005868	39.601	30.998	45. 149	22.672	23. 499	19.410	12.734	17.486
	NT2RP2005876	182.087	242.226	222.167	16.258	31.298	2198, 108	17. 529	20, 489
			63.689	193. 261	46.963	36.817	19.789	29.099	39.512
	NT2RP2005878	91.078					18. 957	6. 938	12.667
30	NT2RP2005883	20. 941	23.594	20.782	9.131	19.950			
30	NT2RP2005886	39. 296	39.439	60.317	47.352	18. 027	22.441	30. 721	46. 169
	NT2RP2005887	57.014	35.877	88.514	16.318	48.626	59.659	24. 351	36. 393
	NT2RP2005890	1.467	3.944	6.429	8.930	1.110	0.000	0. 985	1. 454
	NT2RP2005901	20.981	6.590	21. 187	2.036	7.367	5. 299	7, 158	4, 126
	NT2RP2005902	20.393	16.947	32.820	8.084	22.093	14.130	8. 168	6.766
	NT2RP2005908	151.932	107.992	314.719	54, 159	56. 994	88.516	49. 539	55. 664
35			18.407	16.648	7.455	11, 632	30.787	17.918	15.966
33	NT2RP2005927	44.735						7. 543	21.967
	NT2RP2005933	9.824	12. 141	12.068	9.453	13.104	26.904		
	NT2RP2005941	212.014	56.163	125.056	30.940	64.307	146.736	115. 114	49.381
	NT2RP2005942	18.504	15.139	25.696	8.924	13, 074	17.417	7.750	20.426
	NY2RP2005946	9.728	10.356	21.222	6.005	9.750	8.251	6.713	15, 168
	NT2RP2005970	270.432	161.716	481.318	1122.569	121.562	121.380	132. 328	127, 095
40	NT2RP2005980	46.492	47.170	116.755	26.037	32.671	22.244	18. 314	24.318
	NT2RP2005994	24. 928	29.869	28. 280	11.011	14.761	16, 126	15. 547	12. 085
	NT2RP2006004	33. 199	22.482	40.736	2.254	13. 327	15.670	22.705	28.705
				49.417	14, 196	16.611	24.253	14.883	27.870
	NT2RP2006013	37. 195	30.477						
	NT2RP2006023	352. 327	279.775	760. 112	199.154	108.052	252.378	165. 286	194. 967
	NT2RP2006028	16.154	16.322	9. 466	8. 482	6, 921	16.415	12. 189	19,676
45	NT2RP2006038	0.000	0.000	0.000	2.022	0.000	0.000	2.750	0.000
43	NT2RP2006042	171.799	43.226	84. 802	30.749	34.076	105. 581	87. 203	50, 321
	NT2RP2006043	42.853	34.278	45.615	31.083	20.581	20. 396	21.562	24. 255
			38. 197		22.263	18.783	11.840	20. 855	18.722
	NT2RP2006052	81.736							
	NT2RP2006057	10. 366	16.636		3.253		19.481	5. 521	4. 099
	NT2RP2006064	49.505	48.411	44. 958			35.690	11. 141	42. 302
50	NT2RP2006068	32.753	25. 167	31.742	12.673	13.801	29.984	17.006	20.716
	NT2RP2006069	5.168	1.476				1.811	3. 399	1.150
			28.636			15. 376	32.715	25. 050	58.869
	NT2RP2006071	44. 047							
	NT2RP2006090	36. 345	15. 495				27.073	18.729	16.094
	NT2RP2006092	26.028	24.133			22.737	20.714	23. 958	24.611
	NT2RP2006097	25.828	35.230	63.866	22.123	14. 392	27.780	13.780	24. 430
55									
							•		

Table 95

			A 050 T	- 15 6AC T	1 C46 T	7 606 1	0 507 1	2.890	5.114
	NT2RP2006098	9. 221	8.862	15. 825	1.548	7.695	2.607		
	NT2RP2006099	36.984	26, 268	76.849	17.513	9, 927	22.657	13. 432	24. 422
	NT2RP2006100	6.166	9.812	13. 286	1.403	7.183	10.053	6.143	24. 935
5	NT2RP2006103	61.199	24.990	32.481	5. 365	8.444	14.474	5.643	15. 554
	NT2RP2006106	160, 473	47.046	79.073	14.926	42, 304	95, 141	66.256	54.310
	NT2RP2006127	299.049	72.341	157. 315	35. 299	69.360	160. 904	129, 470	82.790
				14. 868	7. 190	5. 404	8.696	12.032	8.793
	NY2RP2006134	7.925	6.856					16.957	12,817
	NT2RP2006141	34. 209	25.853	25. 279	11.925	12.291	24. 288		
	NT2RP2006166	145.927	143.316	390.446	53.472	49, 950	70.158	31.362	36.423
10	NT2RP2006176	38. 237	32.296	48.672	13.808	41.752	37.097	22.363	19.576
	NT2RP2006181	7.938	2.552	3. 108	2. 599	3.019	2. 533	2.693	7.338
	NT2RP2005184	427,733	164.565	311.744	90.540	136.553	294. 751	209.379	191.687
	NT2RP2006186	9.611	7,571	10.891	2.107	7.906	2.215	13.759	17.231
	NT2RP2006196	64.570	46.625	187.805	24. 294	26.945	31.212	13.067	38.607
			17, 361	28. 888	10.561	7.708	21.719	25, 552	11.042
15	NT2RP2006199	32.521				20.289	14.015	24.697	16.848
15	NT2RP2006200	45.197	30.904	68. 326	12.637				45. 272
	NT2RP2006210	13.063	42.759	41.239	76.812	21.527	21.342	4. 951	
	NT2RP2006219	19.770	12.088	17. 232	4. 165	9. 125	6.702	12.944	14. 193
	NT2RP2006224	56.084	46.968	124. 695	25. 238	22.235	39.796	14.970	39.612
	NT2RP2006237	23.936	13.588	29.768	8. 240_	8.266	24. 478	15.621	12.940
	NT2RP2006238	30.339	10.705	17.681	1.647	9.826	9. 810	5.796	9.385
20	NT2RP2006258	134, 594	65.669	94. 583	35.749	42.774	37.896	67.144	58.117
	NT2RP2006261	30.527	20,607	20.756	7.023	10.500	26.668	20.779	32.986
	NT2RP2006269	273.686	190, 160	282.087	75.118	88.026	221.069	173.956	143.367
	NT2RP2006275	85.280	39.874	56.619	10.486	19, 434	48. 212	55. 210	39.859
		18.372	26.364	78.637	7.247	13.037	10.134	7. 395	10.427
	NT2RP2006282				22.449	24.844	39. 182	9.675	14,948
	NT 2RP2006302	35.243	63. 455	48. 101		27.020	35. 036	41. 172	38. 133
25	MT2RP2006312	65.434	60.394	81.415	26.895				40.879
	NT2RP2006320	42.111_	32.881	107.012	21, 102	25.087	24. 083	19.555	
	NT2RP2006321	7, 504	10. 403	35. 594	9.608	9.770	25. 528	7. 823	3.899
	NT2RP2006323	7.851	2.520	3.223	1.919	1.885	6. 166	3.878	3.640
	NT2RP2006333	30.987	16.865	28.885	6.560	9.086	8. 529	9.411	9. 392
	NT2RP2006334	12.349	6.245	10.111	7,506	2.643	10.779	6.657	9.120
00	NT2RP2006338	3. 452	3.965	5.603	1.571	3.999	1.378	0.000	6.658
30	NT2RP2006339	25.764	16.783	14.506	7.871	9.927	10.052	16.010	8.999
	NT2RP2006355	20.663	13, 101	11.565	6.563	7.455	7. 126	9. 386	6.085
	NT2RP2006365	4. 545	5.794	3.527	6.016	4.317	2. 172	4, 635	2.088
	NT2RP2006374	411.795	181.700	244.772	88.732	81.469	224. 300	186.562	150.290
	NT2AP2006393	49.201	46.271	138.242	24.009	21.170	18.558	17. 331	21, 921
	NT2RP2006394	28. 334	29.547	20.558	4.570	13.741	24.300	15.936	15,737
35	NT2RP2006400			22.520	10.436	6.781	12.164	12.987	14.072
		24. 921	12.448	109.486	76.812	50.885	136.021	80.417	46, 178
	NT2RP2006411	170.083	45.848			17.317	23.371	18.641	17.956
	NT2RP2006429	17.592	22.689	50.747	10.696		37.137	39.774	37.506
	NY2RP2006435	55. 611	34.885	57.426	16.304	26.895		47.063	35.486
	NT2RP2006436	152.017	117.923	294.214	79.789	75.537	107.196		
	NT2RP2006441	24.518	19.297	41.744	27. 285	33.736	14. 991	17. 341	13.076
40	NT2RP2006447	13. 367	6. 103	5.701	2.225	4.629	8. 175	4. 129	2.450
	NT2RP2006454	12. 135	6.375	11.243	2.681	0.000	18.444	3.071	5.464
	NT2RP2006455	11.895	17.452	13.837	6.890	6.   58	14.783	6.071	8.830
	NT2RP2006458	38.021	19. 288	35.373	9.022	12.219	34. 935	12, 195	8.454
	NT2RP2006464	65.475	59, 218	64.107	23.982	11.975	46, 736	45. 415	26.468
	MT2RP2006467	182.556	82, 534	110.746	33.773	58.531	134.845	89. 415	79.911
45	NY2RP2006472	52.035	81.984		27.110	22.246	58.236	23.092	21.013
	NT2RP2006474	87.750	59.508		40, 960	68.884	46.386	41.819	43.544
	NT2RP2006475	31.939	25.175		5. 942	98.476	222.460	20. 356	7.479
		21.072	30.518			6.000	10.383	11.027	21,451
	NT2RP2006476					4. 907	32.045		22.257
	NT2RP2006501	49.705							
	NT2RP2006512	27. 180	32.082				30.488		19.931
50	NT2RP2006526	1.990					0.000		0.000
	NT2RP2006527	89.786	37.810			29.390		41.655	42.817
	NT2RP2006534	12.307	17.082	25.981	7.920	10.780	7.152	5. 503	7.803
	NT28P2006537	152, 141	97, 164				66.047	31.701	24.076
	NT2RP2006543	41.814							15.747
	NT 2RP2006554	5. 859							4.909
55	WITH TOODS	1 3.033	1 3. 31.		7.170	1 0.007			
33									

### Table 96

	NT2RP2006565 8.	67 7.704	24.371	5.814	14. 320	10.696	2.358	4. 111
	NT2RP2006571 279.		116. 541	23.676	53. 970	199. 457	130.143	46, 164
	NT2RP2006573 14.8		14.833	10.165	4. 273	12. 181	5.836	11.189
5	NT2RP2006598 50.		84. 436	36.450	20. 183	47. 448	27.628	33. 428
	NT2RP2006601 363.		103.722	48.729	76. 933	194.071	89.671	34.186
	NT2RP3000002 54.		138. 409	14.410	15.645	42.782	17.893	13.809
	NT2RP3000011 86.		179, 249	26. 157	23, 114	44. 263	20.905	26.577
	NT2RP3000014 13.		34, 145	13.964	62.052	11.790	6.030	23. 999
	NT2RP3000016 37.		44, 744	13. 554	18. 247	35. 947	22. 381	14.827
10			43. 091	11.156	18. 896	66.602	28. 935	18.892
10		342 17.722	80. 534	57. 536	15. 195	28. 526	14. 924	31, 215
			45. 699	14.680	16.043	21.658	37. 591	14. 624
			46.033	9. 722	17. 283	27. 871	22, 418	14. 394
			344. 732	90. 995	100.871	120. 707	93. 233	68. 047
	NT2RP3000037 207.		10. 120	5. 362	4.717	7, 751	13.678	12.858
16	NT2RP3000040 19.		152.312	40.210	22. 300	35.890	26. 992	49. 633
15			156.649	32. 533	24. 374	70.316	23.701	21.537
	MT2RP3000045 66.		49, 113	15. 475	21. 518	33. 173	30.093	27.627
	NT2RP3000047 67.		91, 910	30. 451	29. 572	32.060	28. 583	20. 154
	NT2RP3000049 48. NT2RP3000050 26.		88, 636	24.767	22. 328	23.604	11.688	48. 303
			41.823	15. 685	23. 009	34. 385	30. 860	29.647
	NT2RP3000051 66. NT2RP3000054 102.		100. 267	27.596	31.738	71.470	53.863	44. 388
20	NT2RP3000055 75.		100. 976	32.041	39. 402	46.743	33. 378	38.034
	NT2RP3000056 39.		30. 865	6. 902	18.029	31.675	21. 577	18.143
	NT2RP3000059 37.		41, 439	8.975	11.901	30. 284	16.708	27. 602
	NT2RP3000063 185.		95. 324	25. 648	51.543	102.170	98. 453	32.215
	NY2RP3000068 31.		26. 439	9.761	13. 197	30.638	22. 295	20. 840
	NT2RP3000069 10.		29.064	3. 122	10.074	26.020	12. 191	15. 438
25	NT2RP3000072 14.		11.379	7. 153	9. 559	10.360	3.475	9. 404
	NY2RP3000080 324.		363.840	79.623	88. 104	197.811	132. 385	96.818
	NT2RP3000085 51.		37. 844	10.819	18. 134	39.828	23. 587	17. 525
	NT2RP3000087 17.		46.219	24.865	22. 511	28. 404	15.603	41, 935
		685 15.980	24. 034	8. 335	8. 477	12.472	10.082	10.611
	MT2RP3000109 18.		14, 110	13.437	4. 116	20.790	11.884	10.865
30		508 36.674	39.664	14. 435	17. 248	54, 174	31. 225	36.072
		603 69.403	81.547	42.247	34.639	56.907	38.200	46.977
	NT2RP3000131 120.		90.654	35. 148	31.692	68.253	45.665	51.614
	NT2RP3000134 112.		239. 571	43.058	33.667	26.549	37.483	34. 264
		456 42.787	44. 389	14. 934	21.455	33.205	29.974	26. 136
		473 48.731	52. 053	38.739	20.973	36.445	18.076	21.664
35	MY2RP3000148 63.	507 22.034	36.823	8.026	12.884	44, 451	23. 171	18. 256
	NT2RP3000149 97.	776 30.350	50.788	16.701	25. 676	64.729	43.962	43. 994
	NT2RP3000163 26.	802 19.938	31.411	6.275	9.088	20.951	21.878	23.068
	NT2RP3000168 795.	144 114.786	281. 896	44.650	145. 359	605.075	401.513	129.011
		676 16.941	26. 930	10.017	9.998	22.440	17.412	12.677
	NT2RP3000171 98.	370   112.386	277. 503	71. 994	84. 185	92.446	72.076	90.890
40	NT2RP3000172 61.	369 27.571	34. 375	12.627	22.318	30.658	22.317	17. 859
		000 91.952	162.821	37. 334	35.006	23.969	28.600	28. 365
		123 35.637	164.002	24, 125	21.785	35. 486	29. 445	24. 978
	NY2RP3000201 102.		142, 754	44. 107	29.649	62.714	48.605	33.413
		200 14.164	20.111	7.985	6.611	22.398	6.414	16. 458
	NT2RP3000207 156.		65.015	12.469	27.276	91.928	59.198	23.678
45	NT2RP3000216 198.		109. 849	21.139	46, 927	98. 763	89. 370	46. 993
		042 21.189	35. 304	10.343	13.834	34. 368	22.050	8. 817
		840 11.900		9.467	7.825	20. 185	21.420	5. 118
		369 22.973		25.604	26.475	26.635	21.694	58.778
		604 18.166		7.062	10.046	10.901	14.488	13, 964
		664 54.616		20.000	23.342	34.772	28.379	31.629
50		990 44, 388		18. 177	16.009	48. 324	46.171	18. 108
		735 37.968		18.056	20.915	38. 341	15.352	39.089
		588 21.300		8.851	13. 233	20.777	17.822	10.760
	NT2RP3000251 113	350 59.317	72.549	22.848	36.203	92.438	60.767	66. 928
	NT2RP3000252 60	796 43.494		14.499	17.495	33.029	20. 185	17. 559
	NT2RP3000255 70	857 30.714	38.046	8.728	10.951	.35. 304	33.779	15, 636
55								

Table 97

				00 002 1	10 426 1	13.784	14.416	11,609	16.866
NT2RP3	000262	12.216	24. 325		10. 435	24. 591	38. 289	23. 653	41. 422
NT 2RP3	000266	50.888	57.736		20. 988	9,316	10.177	21. 121	21. 341
NT2RP3	000267	44.661	37.513		10.160	28, 139	32.541	43. 838	26. 269
NT2RP3	000271	83.084	46.052		20.876	21.589	32.926	27.054	36.075
NT2RP3	000278	32.035	43. 936	47.584	38. 351	29.002	49. 528	37. 903	38. 324
NT2RP3	000281	90.519	61.619		27.694		5.631	11.626	1. 198
NT2RP3	000292	3.966	10.376	3. 409	3. 495	2.695	25. 398	24. 814	30.991
	000299	59. 244	17.953	32.272	11.088		68.495	41.001	11. 248
NY2RP3	000304	112.022	42.176	44.039	17. 256	21.312	17.033	12, 288	8. 239
	000310	51.923	40.371	23.866	18.763	12. 225 28. 499	47.636	21.749	17. 055
	3000312	53. 784	42.298	111.962	28.662	34, 370	306.433	171.177	16. 257
		207. 335	105. 256	82.557	32. 315	43.457	53.749	59. 223	53. 805
	3000322	58.959	145.034	68.676	49.667	13.446	25. 783	30, 738	24. 781
	3000324	48.873	14.767	34. 844	16.823	7. 123	38, 932	21.519	21. 276
	3000326	65. 235	51.932	107.139	28.709	30.715	47. 282	30, 786	30.002
NT2RP	3000329	93.768	78.384	210.960	64.677	6. 970	26. 195	18. 445	15. 597
	3000330	24.642	49.689	27.966.	9. 468	1.210	8. 119	6.219	2.641
	3000333	6, 551	4, 474	6.490	3. 373	44. 341	47.850	37.664	24. 434
	3000341	105.554	78.685,	292.105	48.172	14, 171	11.842	9, 663	8. 826
	3000344	21.848	16. 348	18.737	15. 208 4. 341	4, 244	9, 519	7. 200	2. 442
	3000345	13.615	4. 231	8. 891	124.822	216.289	288. 551	206. 453	397. 251
	3000348	215.751	824. 234		26.912	16.935	64. 380	20.038	26. 035
	3000350	75.031	53.082	54. 573 25. 133	36, 113	16.097	67. 120	56, 693	48.617
	3000359	60.599	28.652 40.753	62.678	25. 399	25.559	78.478	40, 608	39.929
NIZKP	3000361	97.227	23. 388	51.997	16.575	24.680	39, 191	19, 302	20. 995
NIZRP	3000366	29, 933	36.646	53. 425	29. 190	18.810	14. 993	29.540	12, 427
	3000378	36. 122 94. 244	64.810	247.061	65. 250	53.993	55. 586	28. 548	35. 998
	3000389	145. 164	130. 566	88.715	60.458	59.767	126.866	45.046	46. 304
	3000393	34, 304	26.482	38.672	12.816	10, 966	53. 247	23. 028	22.722
1012AF	3000395	130.734	261.655	185.074	139.360	67.626	191. 905	113.593	356.673
NIZAT	3000397	23,796	14.400	15, 115	8. 197	10.685	19.437	11.865	15.686
	3000398	53. 315	\$3.724	168, 232	39, 457	13.432	46.057	24. 302	28. 636
	P3000403	57, 006	49, 114	63.081	18.685	24.406	48. 333	25. 226	24. 101
	3000418	50, 531	48. 172	170.356	32.562	26. 123	10.592	29.707	10.504
	P3000424	63. 365	21.340	38.478	16.563	16. 925	53. 214	30.826	17.735
MT2D	P3000427	62.721	39.857	128.557	24.313	28. 283	29.359	22.716	18.652
	P3000431	16.834	8, 211	12. 394	7.692	12.872	8. 065	13.418	13.114
	P3000433	50.616	79.462	104. 236	42.090	39.902	42.064	33. 371	38. 488
	P3000436	16.242	16.422	40.709	15.813	6.539	20.516	10.885	17.733
	P3000439	71.848	23.969	40. 354	14.754	15.239	53.741	31.396	8. 363
	P3000441	11.212	9, 002	12.696	5.044	10.679	13.013	5. 428	5, 597
	P3000444	22.933	18.685	29.664	9.645	13.646	17.025	18. 757	13, 305
NT2R	P3000448	33.060	20.309	55. 374	17.566	24. 368	22.687	16. 155	11.895
	P3000449	6.959	23.459	17. 422	10.472	7.118	8.871	9.364	3.475
	P3000451	125. 446	62.063	59.005	22.337	34. 264	60. 126	41.591	27. 148
NY2R	P3000456	88.916	43.502			29,249	56.726	42.776	13, 869
	P3000460	53. 276	27.765	47. 239	14. 257	14.152	40.035	29.749	40, 267
NT2A	P3000471	120.686	35, 942	65. 409	19.114	21.965	84.016	40.300 64.962	29.801
NT2R	P3000477	135. 254	69.833		56.384	66.460	143.732	14.477	44. 282
	P3000478	29. 313	21.768				33. 287	6.047	2.438
	P3000481	10.750	3.732		0.726	2.539	11.124	24.639	18.179
	P3000484	37.552	21.008		15. 721		24.786 34.015		34.378
NT2F	RP3000487	57.292							10.397
	RP3000512	40.012					78.588	27. 137 42. 509	
	RP3000523	99.365							
	RP3000526	45. 488					36.809		
	RP3000527	44. 308							
	RP3000531	317.473						28, 117	
	RP3000532	69.884							
	RP3000542	53. 226							
	RP3000554	46.760							
	RP3000561	34.700							
NT2	RP3000562	61.916	30.11	37. 119	14.204	13.843	1 33.331		

Table 98

·			16 663	7.228	4.718	6. 130	9.838	13, 311
NT2RP3000578	15. 402	10.162	16.063			17. 246	4.000	23.415
NT2RP3000582	39. 271	21.923	38.385	15.003	10.964			13.885
NT2RP3000584	50, 928	29.642	70.817	14. 592	15. 938	25. 450		
NT2RP3000586	104, 429	33, 153	41.205	15.381	26.618	56.849	60. 938	32.115
NT2RP3000590	26. 385	19, 138	20.258	7.852	11.948	19.961	17.171	14. 281
	38. 458	13. 253	18.849	6.663	11, 214	15.506	10.347	13.365
NT2RP3000592	38.436	124.897	111. 320	54, 127	55, 968	95. 489	58, 183	68.801
NT2RP3000596			21.699	6. 517	10,630	24, 268	12.753	5. 443
NT2RP3000599	27.723	23.836		20.279	17, 695	42. 330	42.704	30. 254
NT2RP3000603	58.661	36.820	44.037			16.695	13.635	14.891
NT2RP3000605	28.480	12.057	23.849	6,629	7.081		13.588	19, 173
NT2RP3000607	24.858	40.289	21.827	5.879	13.852	13.642	6. 395	3. 593
NT2RP3000616	13. 295	18.170	13.744	4. 297	8.368	12.637		
NT2RP3000621	32.066	35, 204	40.136	10.823	13.912	32.917	35.694	30.496
NT2RP3000622	77.250	48. 804	56. 101	26.510	26. 964	60.270	33.756	35.001
NT2RP3000624	69.148	40. 431	50.570	17.495	18.274	44. 392	30.661	19.154
	101. 279	78. 344	315. 194	66.794	47, 806	62.753	39. 571	65.891
NT2RP3000628		57, 931	64.862	38.915	26. 193	49.662	32. 548	66. 985
NT2RP3000631	83. 274	46. 888	137.791	36.803	25. 072	37, 533	39. 161	39.835
NT 2RP3000632	75. 512		37.613	14.925	17,500	33.937	31,430	24.095
NT2RP3000638	42.585	33.637,		71.526	75.904	86.493	86.017	83.257
NT2RP3000644	165. 984	142. 937	393. 193		154.952	264.140	265.679	203.054
NT2RP3000645	406.046	291. 113		137.438	30. 592	38.016	20, 919	70. 560
NT2RP3000652	27. 913	38. 545	66. 305	53.070		41.744	58. 038	42.209
NT2RP3000658	119. 274	49. 302	84. 139	19.097	26.904		54. 638	32.448
MT 2RP3000660	154.015	93.717	291.388	47.970	61.811	77. 178		
NT 2RP3000661	61.960	37.363	58. 907	19.857	23.806	34.888	27. 236	22.377
MT2RP3000665	36.030	11.500	21.945	7.361	8.773	16, 187	15. 502	4. 205
NT2RP3000676	93.465	71.379	82.472	34.775	44. 271	57.208	63.670	56.415
NT2RP3000677	112.363	32.537	52. 925	14.656	38. 145	49.852	47. 252	14. 122
NT 2RP3000681	36.511	66.476	75. 231	35.416	18.401	37.570	41.478	66.253
NT 2RP3000683	58.416	64. 592	97.551	38.537	29.638	37.665	22. 530	57.162
NT 2RP 3000685	114.973	74.466	133. 468	30.843	36.634	44.885	43.642	44. 225
	44.317	22.720	28. 586	11.755	16, 142	19.525	23.913	12. 295
NT2RP3000690	67.409	29, 101	27.424	12.677	18.813	30.558	35. 120	22.330
	69.762	31.242	34. 468	18.438	17, 109	25.677	35.649	27. 340
MT2RP3000708	69. 102	37.708	40. 561	16.843	22. 310	30.132	41.665	29.714
NT2RP3000719	101.619	13. 883	41. 328	20.719	17,808	29.854	31.483	34.754
NT2RP3000721	62. 292		15. 483	9, 343	7.806	5. 356	8, 199	8.869
MT2RP3000728	15. 781	13.248		4.259	5, 390	10.857	12.834	7. 121
NT2RP3000730	16. 503	10.183	12. 261		11.886	24.025	14.564	29.631
NT 2RP3000733	55. 476	33.770	134.994		9, 383	28.210	2.497	11.449
NT2RP3000735	21.669	7.407	9.693		16.809	30.640	25. 306	25. 557
MT2RP3000736	44.789	26.680	38. 153			146, 191	92.971	37. 396
NT2RP3000739	206.032	42.295	130.965		58. 557	190. 392	158. 586	73.831
NT 2RP3000742	348.588	140.895	195. 591		81.126		25.801	41.475
NY 28 P3000753	62.272	31.221	40.211		20.282	94. 033	11.768	12. 157
NT2RP3000759	29.716	22.350			26.712	22.364	14. 289	8.744
NT2RP3000789	39. 203	42.612	22.684	12.737	16.316	24.563		22. 295
NT2RP3000815		54. 520			22.766	48. 540	24. 152	
NT2RP3000818	77.152	41.510				51. 380	29.052	28.761
NT2RP3000820		118. 421	231.975	55. 326		77. 248	35. 255	64. 172
MT2RP3000821	125. 957	64.013		27.822		81.799	57.688	32. 892
NT2RP3000825		0.000		2.088		3.614	3.042	11.736
NY2RP3000826						101.167	61.127	46.725
NT2RP3000836						23. 251	37.753	44. 587
					149, 924	216.645	161.565	295.666
NT2RP3000838			7 657			11.754	5.693	6.807
NT2RP3000839						30. 244	12.610	21.751
NT2RP3000841								28.363
NT2RP300084								48.553
NT2RP300084								22. 982
NT2RP300084								43, 150
NY 2RP300085								8. 905
NT2RP300085								35, 975
NT2RP300085								
NT2RP300086								78.976
NT2RP300086			4 36.13	2 15.94	2 23.416	47. 236	69.109	15. 390
	' - ' '							

Table 99

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	NT2RP3000865	63.270	47.853	102.873	32.472	33. 487	53.656	34. 278	21.893
	NT2RP3000866	34.716	25.903	38, 593	12.760	15,744	54. 423	32. 374	18.694
	NT2RP3000868	85.284	61.512	85.178	31.575	34.644	53. 975	41.313	22. 132
5	NT2RP3000869	77.514	27.048	71.150	21.470	27. 958	26, 061	26.717	11.830
			15.895	28, 790	10.764	12.347	17, 382	19.415	15, 477
	NT2RP3000871	32.339							
	NT2RP30D0875	64.304	26.967	41.187	17.427	17.449	63.004	27.104	29.777
	NT2RP3000895	37,607	26.551	21.094	10. 531	9.611	39.637	23, 121	22.804
				211.235	53.019	47.970	81, 157	50.066	57, 451
	NY2RP3000900	142.017	81.808						
	NT2RP3000901	70.807	27.339	68.215	18.628	38.633	87. 435	34.055	17.677
10	NT2RP3000903	13,003	24. 507	60.511	13.378	13.428	29. 263	6.790	12.691
				31.708	12.964	16.730	32.075	26.793	6. 596
	NT2RP3000904	52.698	18.398						
	NT2RP3000907	166.727	60.470	136.938	38. 479	50.160	105. 219	95. 047	42.673
	NT2RP3000913	94.023	47.327	91.333	23.378	31.301	50. 434	47.912	29. 311
				21.466	16.870	11.875	27, 038	18.723	21.313
	MT2RP3000917	32.888	39.658						
	NT2RP3000919	94.068	33, 556	45. 679	16.703	24. 240	78.449	55. 568	30. 552
15	NT2RP3000921	37.830	26. 534	66. 403	7.357	8.929	61.748	8.623	14.620
						42.178	102.140	75. 932	47.639
	NT2RP3000942	171.953	62.500	108.369	33.025				
	NT2RP3000968	113.182	183.788	251.225	112.172	45. 194	135.391	114. 314	284. 978
	NT2RP3000974	31.061	18.639,	28.044	11.335	13.883	20.765	19.154	8. 182
				144.923	25.869	22.636	53. 158	21, 266	5. 678
	NT2RP3000980	75. 435	43.616						
	NT2RP3000984	80.420	55.909	211.662	30.046	34.753	46.023	41.008	39.028
20	NT2RP3000994	26.597	13, 100	24.899	10.246	18.755	15.021	12.030	11.524
	NT2RP3001001	41.741	14.316	24. 372	9. 822	11.943	20. 519	21.560	7.191
	NT2RP3001004	21.324	19.490	22.465	8.748	12.668	37. 792	8.027	5. 197
	NT2RP3001007	73.322	49.966	175.492	41.711	29.860	30.759	23. 563	18.521
	NT2RP3001012	17.551	14, 673	17. 235	9, 520	7.664	14. 146	11.598	11.610
				40.712	11.357	21.273	42. 340	30, 644	16.851
	NT2RP3001042	56. 542	31, 176						
25	NT2RP3001044	57. 032	39.083	68.934	22.349	40.025	60.364	34. 476	25.794
2,5	NT2RP3001048	39.639	23.540	39.473	18.858	15.279	32. 436	23. 205	26.116
			37.630	102.740	17.755	44.501	73. 595	26.881	21.142
	NT2RP3001050	40.144							
	NT2RP3001055	36. 578	21.787	34.665	11, 391	15, 586	44. 493	17. 343	39.665
	NT2RP3001057	40.477	31.367	56.914	35. 425	16.396	40.782	15. 582	41.540
	NT2RP3001061	35. 545	23.074	31.908	11.906	22, 306	27, 393	25, 460	19.287
							58.467	35. 766	
30	NT2RP3001069	106.748	62.272	150.656	32.917	23. 305			47.515
30	NT2RP3001074	14. 550	14, 541	22.555	7.827	15.140	15. 294	11.052	4.620
	NT2RP3001078	52.226	37.483	. 61.489	16.718	18, 374	26.786	29.722	37.845
			17.926	40.857	14. 999	13.731	23. 258	19. 326	14. 022
	NT2RP3001081	27.544							
	NT2RP3001084	48. 930	20. 162	28.411	8.915	19.688	35. 485	28. 948	20.795
	NT2RP3001095	5. 532	7, 106	9.117	1, 907	1.873	1.686	4. 160	6. 179
	NT2RP3001096	72.786	64.406	72.692	26.305	30. 582	41.52B	32.077	27.965
35									
33	NT2RP3001097	25. 257	17.811	73.704	11.171	12.488	5. 176	10, 401	15. 261
	NT2RP3001107	81.894	32.783	61.356	24. 675	27, 453	53.316	37.116	40.327
	NT2RP3001109	29.099	23.842	24.494	12.892	16, 120	14.893	15. 303	18.912
				36.252	13.681	16.731	44.954	31.601	22.477
	NT2RP3001111	69, 862	29, 991	30.232					
	NT2RP3001112	57.507	80. 536	82.448	80.792	39.380	58. 111	23.819	75.560
	NT2RP3001113	17.615	26.847	19.375	9, 970	8.233	11.421	5.759	13.956
40	NT2RP3001115	21, 858	18.916	28.812	7.324	4, 563	13.477	9. 463	11.057
40								21, 496	
	NT2RP3001116	40.872	22. 335	23.917	10.468	15, 106	15. 973		6.979
	NT2RP3001119	124, 291	38.911	66.173	19.498	29.478	73.564	71.005	23.217
	NT2RP3001120	18.656	32.833	65.009	14.974	1 14, 114	43, 177	17.732	46. 909
		37.515	26.047		9, 469	16.381	17. 926	22.835	12. 549
	NT2RP3001125			38.382					
	NT2RP3001127	11.834	4.025	5. 195	3.694	4.697	2.608	36.686	5.923
45	NT2RP3001133	70. 288	79.857	161.425	34, 123	22.428	47. 525	46,500	34. 323
40	NT2RP3001140	23.850	15, 525	27.441	7.787	14.096	43, 859	22.377	36,073
						1	7- 249		
	NT2RP3001147	41.415	23. 333	25.696	7. 439	15.613	27.307	21.623	12.688
	NT2RP3001150	50.310	27. 305	40.429	13.413	12.407	17. 499	22.391	24. 362
	NT2RP3001152	3.974	1.479	1.712	0.807	0.788	0.915	1.821	0.000
							40. 594		
	NT2RP3001155	19, 961	39.114	41.386	21.748	14, 042		41.468	31.833
50	NT2RP3001156	31.035	17, 102	23.691	6. 973	9, 466	31, 538	17, 411	9.742
50	NT2RP3001159	137. 273	38, 120	74.062	19.455	36.267	73.862	75.135	35.944
						20, 302	35, 625	22.021	14. 394
	NT2RP3001170	35.615	34. 235	64.722					
	NT2RP3001175	58. 889	60.413	127.466	30. 928	33.027	62.693	23, 996	56.392
	NT2RP3001195	72.627	47.832	119.011	16. 902	19.658	15, 312	25.740	27.006
	NT2RP3001209	458. 437	263.607	330.947		187.739	350. 320	327.764	223.342
	MI CVL 200150A	1 430. 43/	1 203. 601	1 330. 341	1144. 032	1 101.133	1 330. 320	1 451.104	1 220. 376

### Table 100

					40 700	3A 1A2 I	2 4/2 1		12 200
	NT2RP3001214	15.750	24. 578	18.804	10.536	12.107	7.011	7.277	12.208
	MT2RP3001216	29. 191	41.925	42.777	28.031	31.602	30.104	23. 133	26. 408
	NT2RP3001221	25. 240	20. 176	20,644	4.858	10.770	7.315	13, 513	8. 946
5	NT2RP3001226	54. 304	47. 592	67.642	24. 341	24. 384	38, 331	32.669	45. 697
-				27.561	11.735	11.032	25. 537	13. 367	12, 758
	NT2RP3001230	23.680	20.599			6.746		5. 130	7.513
	NT2RP3001232	4. 151	14. 071	13, 135	9.855		9.778		
	NT2RP3001236	28. 593	14. 443	35, 687	7.512	5. 884	10.789	13. 692	14. 135
•	NT2RP3001239	15, 380	5.089	16.960	4. 419	3.497	6.366	6. 599	8. 021
	NT2RP3001240	17. 531	13.481	30, 743	12.073	14.733	12.342	22. 647	11.311
10	NT2RP3001245	17. 405	11. 231	97, 349	10.570	10. 567	11.712	11.709	5. 360
•	WT2RP3001253	29. 416	21. 939	10. 308	17. 125	8.767	28.879	20. 229	15. 732
			26.700	30, 561	28. 122	16.780	50.988	35, 111	9. 149
	NT2RP3001259	66. 464				11. 179	5, 131	7.611	6. 513
	NT2RP3001260	15.811	4, 776	8.508	3.773		13, 645	11.466	13. 410
	NT2RP3001264	17. 474	9, 326	19, 891	6.147	0.000			
	NT2RP3001268	10.917	11, 531	28. 253	21.540	5. 251	19.724	6.886	19.730
15	NT2RP3001271	504. 472	230.117	363. 954	129.052	147. 454	341. 938	290.090	221. 262
	NT2RP3001272	53.274	65. 558	170, 406	46.512	25.065	49.775	25. 165	40.714
	NT2RP3001274	379.452	180.634	305, 168	109.916	102.975	251.219	221.619	125. 753
			44. 463,	39. 465	12.598	20.694	29.868	27.346	15.608
	NT2RP3001275	69.350		58.661	22.588	41. 962	35. 792	31.005	24.067
	NT2RP3001280	84. 373	66. 148					37. 860	23, 491
	NT2RP3001281	108.112	65.094	147.713	26. 972	40.778	44. 735		57, 051
20	NT2RP3001288	37.247	71.613	48.891	21.593	34.714	52. 211	28.610	
	NT2RP3001297	74.827	48.767	64.501	33.081	24.851	55. 981	. 45, 160	38. 893
	NT2RP3001300	97. 287	54, 906	120.465	40.784	42.784	70.008	44. 204	40. 196
	NT2RP3001301	11.093	5. 654	18.227	4. 517	6.710	15.021	4. 763	1.496
	MT2RP3001307	61.481	16.300	67. 269	13.678	18. 372	43.312	26. 354	11.694
	NT2RP3001310	25. 947	50, 116	44, 928	47.745	27.742	21, 494	12. 532	23. 321
	NT2RP3001318	2.615	3.369	14.422	2.026	3. 965	19, 407	1.976	0.653
25	M12RP3001322	23, 311	16, 139	27.515	12.075	11.630	21, 825	11.372	27. 956
	NT2RP3001325	22.066	21. 492	31.828	22.944	8. 193	35.016	13, 134	12.705
		267.619	127. 929	200. 245	81.462	81.219	191.701	150,006	129.805
	NT2RP3001338	55. 924	18. 296	23.218	9.542	15.077	30.484	15. 924	10. 368
	NT2RP3001339			242.840	118.851	106. 391	255, 313	197. 733	160.604
	NT2RP3001340	298. 177	147.842			14.654		20.669	6. 157
••	NT2RP3001341	23.654	19.357	26,001	10.758		8.713		
30	NT2RP3001354	87.315	79.863	264.818	54.210	48. 577	53.865	34. 407	62. 241
	NT2RP3001355	42.549	24. 220	47.797	11.284	26.805	23, 247	21.876	15. 122
	NT2RP3001356	34.895	26. 366	50.692	16.458	11.954	15. 544	17.696	14.918
	NT2RP3001359	69.545	40.643	64.520	10.543	19.486	38.410	36. 229	16.040
	NT2RP3001364	52. 551	18, 103	37.863	13.181	12.916	34, 493	28.810	10.544
	NT2RP3001373	92.853	21. 226	85. 327	12, 110	28.221	75.073	40.142	19.775
35	NT2RP3001374	18.567	16, 153	13.874	9.085	8.007	19.729	12.896	13. 227
55	NT2RP3001383	35.886	15,749	32.731	12.969	13. 335	20.056	21.243	6. 300
	NT2RP3001384	48.057	31, 309	50.523	17.718	21.014	25.468	27.812	18. 039
	NT2RP3001388	55.759	50. 699	117, 391	21.210	51, 970	52. 288	26.887	29. 189
			21. 933	25.705	8.332	7. 588	8. 588	10.071	8.795
	NY2RP3001392	21.410	8, 348	19, 141	7.594	10.677	11.741	7. 988	10. 281
	NY2RP3001396	15. 219				59. 027	175. 527	102.665	51.488
40	NT2RP3001398	232.068	78.008	227. 136	38.826 33.588	29. 343	44. 058	36. 467	18. D64
	NT2RP3001399	92.466	61.566	193.463				12. 991	23. 925
	NT2RP3001402	26.552	22.030	30.054	10.244	16.784	15. 120		
	NT2RP3001407	18.523	26.250	28.873	22.708	11.616	32.784	12.119	9. 034
	NT2RP3001416	45.040	28.810	36, 947	10.094	15.710	31.887	29, 218	27.952
	NY2RP3001420	39, 104	40.226	112.497	29.782	17. 944	17.648	21.378	24.813
	NT2RP3001425	39.881	24. 233	38. 220	17. 938	17. 233	24.809	27. 932	23. 131
45	NT2RP3001426	93.587	58, 250	74, 483	29. 242	37.668	70.217	46.545	43.211
	NT2RP3001427	42.182	35. 278	33. 424		17.612	29.923	26.039	24.764
	NT2RP3001428	59.474	65, 787	162.966		19, 191	22.870	32.859	22. 870
						8, 727	11.508	11.358	8.026
	MT2RP3001429	35, 365	23, 903			8.729	9.061	7.315	14.038
	NT2RP3001432	42.083	23. 762						
50	NT2RP3001439	136.789	39.813			30, 564	95. 253	70.005	58.770
50	NT2RP3001441	38.061	24.064			14.885	31, 263	17.597	19.725
	NT2RP3001446	20.584	15.857	21.782		7. 326	13.920	17.301	7.485
	NT2RP3001447	104.606	70.020	154, 062	30.513	35.791	54. 330	36.473	40. 409
	NT2RP3001449	10.642	11.657			3.069	9.786	3.931	5.883
	NT2RP3001453		43.788			14.076	36. 585		35.481
	MI CKF 300 1453	87.023	1 74. 180	1	1 22. 500	1 , 1, 0, 0	1 - 3, 434	,	

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Table 101

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				40 475	0.474	16.537	32.376	23.383	23.793
	NT2RP3001457	57.656	31.667	38. 475	9.474				13.216
	NT2RP3001459	60. 291	21.305	34. 270	9. 400	12.047	30. 246	18, 427	
	NT2RP3001463	37.349	24, 189	26.737	11. 241	16.712	12.719	16. 251	18.600
5	NT2RP3001466	3.829	2.179	4. 207	1.152	6.985	7.668	4. 907	8.467
	NTZRP3001472	42.523	90. 955	71. 226	30.689	20.551	29. 208	32.709	50.536
		78.059	60.351	58, 086	17.203	25. 592	46. 882	39. 257	36.566
	NY2RP3001475		39, 412	55.653	11.108	26.361	52. 488	31.590	18.401
	NY2RP3001479	51. 578		39, 150	6.364	17.825	19.656	7.865	7.287
	NT2RP3001490	9.839	19.316			12.384	24.009	18.581	38.062
	NT2RP3001492	26.968	22.905	24. 652	26.603			27. 157	19.314
10	NT2RP3001495	42.340	19.294	36.741	7. 565	17.241	28. 985		
	NT2RP3001497	32.950	17.434	21.044	7.024	15.546	10.180	19. 393	11.452
	NT2RP3001501	49.067	12.638	47, 469	8.720	17.879	41.926	36.474	34. 151
	NT2RP3001527	128.120	105.243	244.961	55.672	47.467	62.628	70.008	82.431
	NT2RP3001529	126. 912	81.307	206.759	35.211	39. 398	67, 609	39. 145	62.778
		88.926	38, 255	69.884	13. 233	24.804	68.411	33. 275	12.991
	NT2RP3001538		43, 540	51.302	22.808	20.905	63.546	26220	38. 541
15	NT2RP3001539	81.817		19.344	6. 489	4. 478	17.599	4,710	7.688
	NT2RP3001542	11.704	7.892		28. 542	25. 159	35, 315	25.069	26.210
	NT2RP3001549	60.840	55. 102	62.218		24. 143	31.920	38.546	16.779
	NT2RP3001554	63.142	38.335	57.520	12.016	2.833	51.650	11.927	4.890
	NT2RP3001560	31.508	10.439	17, 431	4, 171		73.893	63. 557	42.901
	NT2RP3001561	63.493	90.177	97.829	34.619	16. 230			54. 688
20	NY2RP3001564	24.224	31.924	65.851	31.318	22.874	32. 192	22.750	
20	NT2RP3001568	67.785	39.398	77.618	15.998	21. 374	60.561	47.360	27. 334
	NT2RP3001575	158.363	105.187	188.761	35. 371	49. 236	104.929	66.520	52. 127
	NT2RP3001580	22.928	24, 103	27. 902	11.308	13.846	10.773	15. 209	6.535
	NT2RP3001587	30.882	46.805	32.389	23.716	21.127	18, 550	19.430	26.668
	NT2RP3001589	87. 238	55. 913	140.234	21.405	30.269	16.502	28. 129	10.227
	NY2RP3001592	47. 242	30.596	31.040	13.899	18. 557	41.892	35, 638	42.607
. 25		16. 545	13.286	20.677	4, 980	9.882	24.464	11.354	5.914
	NT2RP3001607		35.856	58.646	18. 572	27.828	41,340	38. 549	32.556
	NT2RP3001608	107. 899		94. 058	22.958	35, 402	79.493	85.697	41, 703
	NTZRP3001613	181.447	52.790	28. 424	19.581	14.720	20.892	19, 236	19, 461
	NT2RP3001619	37.170	25.761			14.008	11.806	23.506	15.754
	NT2RP3001621	25.051	25, 597	20.759	20. 248	14. 221	12.517	24.496	15.072
	NT2RP3001629	42.495	29.023	21.485	11.692		15.299	24.396	22. 471
30	NT2RP3001630	55. 203	33.318	12.380	8. 398	13.075		12.424	22. 548
	NT2RP3001631	44. 095	28. 385	25.774	21.960	8. 104	12.247		25. 097
	NT2RP3001634	49.389	31.519	50. 276	17. 438	9. 120	14.725	16.971	
	NT2RP3001642	58.384	63.135	64. 537	32. 197	35.654	40.765	40.711	48. 812
	NT2RP3001646	46. 102	25.499	30.071	11.012	13.561	30.364	19.040	15. 478
	NT2RP3001650	24.560	13.692	28. 286	3. 177	10. 587	18.321	16.939	9. 216
35	NT2RP3001667	25. 379	40.979	30.064	11.709	14, 158	32.432	17. 482	25.227
55	NT2RP3001671	51.796	35.962	30.710	14.900	16.883	48.652	22.108	17.635
	NT2RP3001672	125. 298	47.766	73.324		41.587	103.311	68. 493	24.949
	NT2RP3001676	44. 058	36.932	114.623		23.379	25.887	17. 997	8.670
		48. 527	41,805			18.855	29.685	32.419	36.221
	NT2RP3001678	56. 508	36.021	81.826		18.731	40, 182	31.070	17.889
	NT2RP3001679		20. 214			10.046	16.063	14.268	5.567
40	NT2RP3001682	33. 136 95. 365	62.809			21.045	38, 439	16.225	11.304
	NT2RP3001685					41.328	31.580	55.067	32.257
	NT2RP3001688	122.935	103. 280			17.447	39, 119		18.809
	NT2RP3001590	48. 596	45. 935			26. 960	72.114		19.480
	NY2RP3001693	76.315	27.860			60. 841	9, 615		9.560
	NT2RP3001696	35.875	28. 246			27. 452	36.516	25.269	42.349
	NT2RP3001698	43.726	102.017				11. 104		20.780
45	NT2RP3001708	36.121	26.604	23. 16	16.082	1.714			61.807
	NT2RP3001712	113.609	129.822	366.56	5 126.311				
	NT2RP3001716	9.845		13.73					
	NT2RP3001724	43. 121	_	32.82	0   19,574				
	NT2RP3001727	72.718					61.017	47. 265	
	NT2RP3001729	10.639						64.178	7.190
50									
	NT2RP3001730								
	NT2RP3001733	40.642							
	NT2RP3001737	106.767							
	NT2RP3001738								
	NT2RP3001739	119.404	43. 83	7 72.50	1 19.33	38.072	77.99	61.245	67.161

Table 102

		- CO 301 T	20 572 1	86.234	43, 100	39.678	62, 316	23.594	32.745
	NT2RP3001742	58. 731	59.672			31, 536	36. 350	15. 916	. 18. 927
	NT2RP3001751	48. 631	34.876	158.212	47.830				38. 702
	NT2RP3001752	94. 578	61.575	307.338	43.572	55. 894	46. 187	9. 168	7. 574
5	NT2RP3001753	23. 594	18. 268	28.874	16, 113	17. 103	13. 403	14. 360	
	NT2RP3001754	257.019	147.414	145. 593	48.124	69.378	138.023	89. 833	70.678
	MT2RP3001756	106. 542	23.060	11.890	3.761	12.461	39. 172	8. 157	5. 587
	NT2RP3001764	97.616	41.097	57.216	18.829	29.263	46.634	32.748	8.673
	NT2RP3001771	89,626	20.149	49.519	15.739	25.796	66.030	41.963	10.077
	NT2RP3001777	58.067	26.504	49.752	19.057	29.401	31. 279	31, 451	13.675
10	NT2RP3001782	78. 349	53.349	189.787	42.036	31.814	40.007	32. 537	31.265
	NT2RP3001792	116, 784	33.273	79.271	30.838	34.190	79.914	66.384	24. 845
	MT2RP3001799	56.002	33.221	58.797	25, 754	25.042	47.831	44.737	16. 237
	NT2RP3001819	99. 523	31.676	64.535	11,784	27.979	48. 855	30.729	15. 920
		73. 466	107.350	119.232	72.609	47.731	75.897	53.911	85. 472
	NT2RP3001829		27. 404	43.716	32.034	20.484	30. 135	10.824	26. 221
	MT2RP3001836	24.805	48. 291	49.763	22, 383	28. 432	53. 489	36.072	27. 184
15	NT2RP3001839	65.164		123.313	25. 118	28.657	41.010	27, 431	29. 936
	MT2RP3001844	66.622	61.308		46.040	30.799	64.847	88.349	81.167
	NT2RP3001848	155. 399	71.963	136.546	25.627	11.291	39. 721	17.078	15.781
	NT2RP3001854	27.874	31.416,	19.202		8, 116	5. 497	12.706	16.492
	NT2RP3001855	27.658	6. 272	33.869	13.508	19, 378	31, 136	31.027	10. 998
	NT2RP3001857	56.318	28.077	35, 198	13.759	15,411	32. 167	36.372	11.561
20	NT2RP3001858	54. 103	24, 171	29.092	13. 284		45, 119	47.585	13.999
	NT2RP3001861	63.497	29.741	57.635	20.968	28.106		11.074	7. 998
	NT2RP3001866	10. 249	12. 382	19.920	12.616	11.772	42. 526		4.517
	NT2RP3001871	12.631	15.883	25.471	6.868	6. 207	12.620	4.571	
	NT2RP3001874	11.507	11.103	18.203	4.856	8.061	6. 546	18.725	3. 916
	NT2RP3001878	18.465	9. 045	11.792	9. 332	8, 403	9. 161	9.699	4. 707
05	NT2RP3001885	96.791	37.635	150.137	59. 749	39. 678	65. 282	51.265	28. 873
25	NT2RP3001896	32.191	20.738	27.405	6.654	24, 453	44. 306	22.893	9. 765
	MT2RP3001898	78.914	42.917	61.453	15. 826	29. 295	67.204	51.298	17. 212
	NT2RP3001899	41.343	15. 205	21.780	9, 260	12.053	26.711	26.329	25.656
	NT2RP3001901	66.535	31.714	47.183	21.483	19.792	40.418	25.763	53.079
	NT2RP3001915	13.485	9. 383	12.294	10.822	7.631	16.078	5, 131	7.213
	NT2RP3001926	6. 261	3.066	9. 593	3.684	3.576	9, 671	11.215	1. 684
30	NT2RP3001929	60.492	34.768	142. 251	36, 157	39.929	21.055	30. 245	40.792
	NT2RP3001931	61,641	53.696	67.258	14, 577	19.384	29. 503	29.562	27.881
	NT2RP3001938	40.274	25. 723	28.062	7.496	13.890	31.768	21. 367	10.885
	NT2RP3001943	28.287	39.405	55. 585	15. 302	25.639	35.454	26.626	14. 424
	NT2RP3001944	73.315	27.407	47.229	18, 622	23.648	23. 459	28.532	14.827
	NT2RP3001945	34.740	226.973	44.000	46. 158	19.151	46.315	28.688	17.572
35	NT2RP3001947	116.378	37.593	58. \$70	24, 995	34.634	68, 127	58.533	46.304
**	NT2RP3001949	21.954	11.535	33.877	4.860	16.683	22.117	14.558	17.598
	NY2RP3001952	143, 519	121.088	53.648	50.889	37.440	105.617	83, 380	63. 243
	NT2RP3001954	62.996	26.992	48.377	12.537	20.542	32.191	29.976	25. 668
	NT2RP3001956	129, 978	158. 142	151.322	123. 162	62.713	92.406	67.282	100.024
	NT2RP3001967	93.636	55.466	88.272	10.572	29.097	36.626	46.055	17.092
	NT2RP3001969	34. 479	21.534	19.898	9, 167	5. 399	15, 105	15.158	2.531
40	NT2RP3001975	37. 230	23.786	60.518	23.795	22.136	24.440	19.911	25. 309
	NT2RP3001986	24.216	19.727	27.547	10.801	12.852	13.805	18. 920	10.726
	NT2RP3001989	1.471	1. 909	7. 536	0.621	1.861	0.578	0.269	1.159
	NT2RP3002002	86. 258	90.727	227.536	60,750	55. 252	43.279	35.951	27.250
	NT2RP3002004	19.703	13.852	27.972	4.752	16.286	18.094	19.787	7.343
	NT2RP3002007	23.474	20.861	30.066	11.557	12. 246	16.556	11.639	9. 539
45	NT2RP3002014	73.272	44.064	105.038	21.583	22.923	30.079	37.416	19, 158
		45.650	25. 353			11.588	23, 493	22, 893	14, 440
	NT2RP3002015					2. 555	1. 242	5. 234	1.639
	NT2RP3002033	7.919				3. 123	8. 022	6.419	4. 266
	NT2RP3002045	21.618					15. 344	8.332	4.770
	NT2RP3002054	12.875							
50	NT2RP3002056	15. 165					12.472	5. 599	27. 199
50	NT2RP3002057	34.454					23.460	21.618	18.390
	NT2RP3002061	35.549	24.492	34.009	18, 402		21.477	15.115	17.613
	NT2RP3002062	30.631		52. 221	11,461			8.319	7. 954
	NT2RP3002063	23.330				12.276	13. 149	10.874	7.143
	NT2RP3002064						47.256	44.374	26.732
	MI TYL 2005004	100. 543			1 11111				

Table 103

	TUTA 504 504 504 50 73	10 641	8.678	10.550	3.877	8.890	15.118	11.681	9.986
	NT2RP3002071	18.641		17. 244	8.390	7.984	14.893	9. 734	12.810
	NT2RP3002073	21.421	28. 270						19. 857
	NT2RP3002074	58. 380	28. 105	42.899	18.734	20.881	18.721	29.611	
5	NT2RP3002075	59. 306	37. 344	42.700	25.078	27.978	35. 950	13. 241	20. 022
	NT2RP3002877	120.301	28.839	29.039	10.364	15.319	40.212	29. 213	9.478
	NT2RP3002081	26.831	15.778	21.982	12.572	10.820	14.083	12.614	11,083
	NT2RP3002086	87.926	53.777	142.446	48.023	26.542	32.148	26. 246	52.677
		33.062	35. 549	57.575	42.152	21.321	27. 515	18.554	12. 485
	HT2RP3002094		22.516	39.610	9. 388	18,743	33, 193	33.700	12.017
	NT2RP3002096	49.540			21.007	14, 939	24. 394	20. 920	11.430
10	NT2RP3002097	26.334	27.838	34.989				24. 025	31.657
	NT2RP3002098	44.592	23.806	37.622	21.688	21.108	22.573		
	NT2RP3002102	79.033	86.261	164.477	46. 235	35. 252	56.190	33. 162	43.258
	NT2RP3002106	77.525	71.059	239. 471	34. 504	16.297	23.309	16.557	12. 205
	NT2RP3002108	44.613	18.028	23, 167	12.003	9, 700	17.108	11.361	8.970
	NT2RP3002109	48, 832	54. 217	110.537	30. 507	53.885	32.217	28.672	32.057
	NT2RP3002110	89.630	210.042		193.998	55. 568	79.385	66.216	96. 572
15			35.313	57. 256	20.790	24.151	40.633	31.916	21.890
	NT2RP3002113	56.372			14. 039	12.431	13.596	15. 152	8. 244
	NT2RP3002120	29. 242	37.086	18. 529				15.462	9, 151
	NT2RP3002121	16.794	22.468,	34. 546	15. 934	19.042	18.137		
	NT2RP3002126	41.432	79.714	33, 116	16.398	35.960	52.883	34.750	31.846
	NT2RP3002128	181.295	79. 422	107. 432	30. 207	38.340	110.226	72.274	55.110
00	NT2RP3002130	146.473	43. 354	77. 922	29. 452	37.242	74.976	38.796	20. 167
20	NT2RP3002133	57.753	91.578	70.347 -	18.863	21.214	49. 924	14. 482	21.057
	HT2RP3002136	43.801	49. 959	66.820	35, 859	53, 999	51.027	15.709	17.711
	NT2RP3002140	64. 973	38. 168	59.056	29.445	31.803	46. 421	49. 899	13. 225
			135. 567	308. 150	95.713	104, 450	105, 460	76. 193	111, 169
	NT2RP3002142	132.430				54. 554	46. 952	38,770	22.003
	NT2RP3002146_	110.073	59. 842	274. 145	50.104			36. 864	20. 569
25	NT2RP3002147	79.974	78, 251	76. 290	23, 131	32.938	49.028		
23	NT2RP3002151	28. 317	56.044	35.024	31. 238	13.466	19.730	29. 531	21.213
	NT2RP3002155	113, 358	59.837	83.053	31.667	33.044	85.787	59.718	11.547
	NT2RP3002156	18. 567	17, 466	43.089	11.697	14, 283	20.150	19. 476	8. 599
	NT2RP3002160	45. 470	32.287	51.148	8. 537	17.337	18.576	19.383	9. 987
	NT2RP3002163	58.319	76. 385	85. 220	36.452	25, 979	54. 323	41.118	65.634
	NT2RP3002165	99.653	52.118	87.449	32.574	44. 305	65.099	54. 567	25. 366
30	NT 2RP 3002 166	37, 449	18.398	38. 523	7.973	18,270	16.300	16.573	5.836
		138. 293	67. 332	233. 564	25. 504	39, 519	46. 406	22. 234	32. 147
	NT2RP3002173		25. 592	20. 612	10. 322	10.075	33.100	18. 166	8. 352
	NT2RP3002174	34. 983			15. 521	6.186	13.861	17.883	5. 289
	NT2RP3002181	25. 553	17. 452	12.477				23. 439	7.852
	NT2RP3002185	130.901	22. 501	42.897	20.805	18.996	58.093		
	NT2RP3002193	48.914	35. 893	57. 402	12.166	28.331	65, 610	51.617	21.157
35	MT2RP3002204	25. 437	16.825	30.602	6. 124	18.001	26. 166	12. 479	21.873
	NT2RP3002244	49.842	27. 141	57. 904	22.937	24.682	26.606	32.340	22. 561
	NT2RP3002248	86.580	63.454	102.977	40.434	37.198	51.108	39, 002	34.672
	NT2RP3002253	55. 575	9, 382	8.780	13.506	12.566	16.080	15. 217	3. 963
	NT 2RP3002255	35.015	68.339	52.684	56.744	24.356	32.145	25.739	37.424
	NT2RP3002284	55. 986	34.735	59.125	27.856	28.745	42.746	33.939	8.983
	NT2RP3002267	80.099	23, 461	44.639	24. 189	20.404	52. 393	26. 915	33. 436
40	NT2RP3002273	112.221	85.604	140.868	66, 160	58.014	79, 427	50, 417	36.059
				50. 683	13.361	24.974	43.308	34. 452	31.732
	NT2RP3002276	62.303	48.041			13.790	21.545	20. 931	8.966
	NT2RP3002281	40. 333	19.037	24. 587	16.378			13. 138	14.040
	NT2RP3002286	27. 525	24.696	32, 519	15.907	12.207	12. 167		
	NT2RP3002297	184. 330	104.754	239. 133	101. 492	75.626	106, 831	74, 738	83.240
45	NT2RP3002301	53.311	19. 361	38. 416	18.640	28.458	40.874	31, 521	16. 259
45	NT2RP3002303	151. 906	66. 595	108.440	41.097	41.354	98. 439	62.889	20.317
	HT2RP3002304	9.712	7.368	13.268	9.520	3.566	6.387	8. 272	2.623
	MT2RP3002309	34.656	9, 379		19.687	8.915	31.244	28,005	8.625
	NT2RP3002311		21.425		9.614	15.336	23.060	17, 155	24.047
	NT2RP3002315	44. 224			29. 239	27.551	69. 218	44. 550	30.664
		60.149	39.087			8.358	20. 152	26, 375	28.658
50	NT2RP3002319	29, 909	14. 381				55. 982	49. 196	49. 374
	NT2RP3002324	84.644	48.794		26.759	38.717			24. 194
	NT2RP3002330	40. 225	35.781		18.069	24. 353	43. 432	29.047	
	NT2RP3002333	739.604	109.838		63. 516	145.604	638. 213	358. 164	89.849
	NT2RP3002337	12.429	9.488				6.399	6: 548	8. 159
	NT2RP3002342	18.485	16.965	24.764	8. 272	19.656	13.221	7.806	10, 971
								_	

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### Table 104

/ <u>-</u>					44.	37.461	51.737	39.000	34.837
	HT2RP3002343	98.077			34. 235				11.298
1	NT2RP3002351	11.568	8.544	17.447	8.504	7.516	10.032	16.378	
	NT2RP3002352	61.768	50.393		25. 295	17. 190	34. 146	31.668	29.346
	NT2RP3002353	84.753	66.818	124.498	39. 521	45.715	83. 255	42.335	39.394
	NT2RP3002362	147.017	77.918	101.793	33.659	48. 293	105.808	93.191	47.902
	NT2RP3002363	51.360	22.194		16.354	18.149	41. 241	27.368	9. 958
		22. 585	15. 479		11.831	11.702	22.164	19.250	14.688
	NT2RP3002377		26, 590		12.961	18.317	29. 595	32, 435	19. 372
	NT2RP3002383	36.652			16. 223	19.758	13.702	16.544	34, 308
	NT2RP3002388	41.759	29, 432			23, 109	44. 424	35. 200	24.054
	NT2RP3002394	64.877	31.565		18. 641			334.003	155. 648
	NT2RP3002398	344.708	216.589		53. 561		244. 214		92.415
	NT2RP3002399	120.898	118.841		92. 322	61.939	76.458	34. 837	
	NT2RP3002402	52.959	35.232	68.571	16.571	20.492	53. 151	21. 545	24.518
	NT2RP3002409	167.688	37, 697	100.184	25.069	35.882	114.827	88. 945	40.800
	NT2RP3002410	144. 081	109. 377	101.178	45. 575	39. 226	71.259	45. 433	41.401
	NT2RP3002411	93. 030	33.468	50.254	10.997	27.600	27. 023	23.738	15.047
		43, 781	19.997	33.403	9.720	14, 797	31.472	21.609	8.498
	NT2RP3002429		12.378	25.831	8.000	12.388	14. 483	16.180	11.704
	NT2RP3002448	18.505			11.518	12.679	23.830	18.696	7.724
	NT2RP3002454	22. 834	27.433,	27. 109		25. 184	40. 843	26. 300	25.891
	NT2RP3002455	42. 267	39.024	48. 252	18.078			34, 943	107. 915
	NT2RP3002456	63.618	62.895	132.023	60.865	48.457	47. 502		23.685
	NT2RP3002462	81.232	66.732	75. 545	22.706	28.463	63.509	41.976	
	NT2RP3002469	31.281	25.018	41.900	16.283	18.312	31. 313	22.887	8.884
	NT2RP3002470	394. 179	240.381	344.971	150. 134	156.904	226. 629	242.639	129.974
	NT2RP3002484	119.962	120.572	179.767	55. 590	78.186	80.561	80. 333	27.126
	NT2RP3002491	20.237	11.861	12.690	4. 514	6.231	7.954	11.431	9. 537
	NT28P3002494	102.258	227.475	73.714	31.409	28.100	91, 250	58. 572	81.116
		111. 163	46.894	64.415	16.949	25.888	63.935	42.893	24.093 .
	NT2RP3002497	77.111	26. 529	42.337	12.959	16.485	30, 996	37.915	22. 524
	NT2RP3002500		44. 526	44.009	16. 212	22.884	27. 120	37. 461	16.746
	NT2RP3002501	53.661		40,061	20.054	21.830	23. 291	29.988	18. 925
	NT2RP3002512	63,608	44.357			22.514	23. 399	23. 938	31.672
	NT2RP3002529	45. 341	43, 112	48. 262	25. 498		49. 543	39.779	31.520
	NT2RP3002533	94. 195	65.870	61.041	18.300	73.412		29. 822	42.332
	NT2RP3002539	48. 864	37.046	54. 572	30. 194	21.685	26.897		11.502
	NT2RP3002540	30.794	21.358	37. 383	11. 560	13.724	17.298	19. 581	
	NT2RP3002543	223.940	110.144	120.839	52. 219	64. 994	144.657	115. 227	76.872
	NT2RP3002545	15.100	41.894	32.270	19.423	32.049	13. 151	11.195	10.417
	NT2RP3002549	28.199	14.150	27.495	13.528	19.671	17. 420	11.163	7.548
	NT2RP3002552	47.064	17, 945	25. 504	12.370	13.372	28. 220	22.837	14.570
	NT2RP3002558	61.923	30,846	56.966	17.185	28.359	33. 407	22.300	21.755
	NT2RP3002565	62.350	42, 196	107.270	25.722	27.937	33. 279	27.380	20. 262
	NT2RP3002566	54. 275	39.776	49.593	22.587	24.849	18.616	38.067	25.776
		16.476	11.788	20. 308	3.165	5.305	12.738	11.591	7.492
	NT2RP3002571	65.635	36, 206	37.772	17.526	23.615	29.016	17. 205	16.571
	NT2RP3002572			49. 387	56.147	11, 324	27.549	32.818	43.821
	NT2RP3002573	104.009	83. 178	33.591	12.529	6. 690	22.718	19.368	7.491
	NT2RP3002577	52.884	22. 337		21.690	15. 920	21.971	21.241	10.888
	NT2RP3002579	71.729	30. 291	35.007		41, 904	56.964	46. 155	37.227
	NT2RP3002582	81.979	51.167	67.043	31.231	19, 982	21.677		12.145
	NT2RP3002587	26.087	32.407	69. 922	18.487				4.548
	NT2RP3002590	7.512	8. 105	10.729	21.190		8.973		9.770
	NT2RP3002602	47.775	17.298	29.784	12. 271	15.119	25. 375	31.820	
	NT2RP3002603	161.708	183.767	216.650	65.839		109.597		115.706
	NT2RP3002521	119.248	24.598	40. 553	16. 479		62.060		25. 390
	NT2RP3002622	69,767	50,020	145.390	29.140		41.045		15. 918
	NT2RP3002624	1.393		0.000	0.942				1,562
	NT2RP3002628	9, 999		17.715			14. 530		5. 659
	NT2RP3002629	249.675		98. 304	56.623		134. 353	115, 158	40.132
					0.000				0.000
	NT2RP3002631	0.595		27.336			15.918		14.888
	NT2RP3002647	30.462					17. 229		30.637
	NT2RP3002649	120. 351							51, 110
	MT2RP3002650	78. 123		55. 575					10.676
	NT2RP3002652	40.736							1
	NT2RP3002654								
	NT2RP3002657	79.710	86.415	129. 177	41.769	103.657	80.846	59, 737	46. 192
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Table 105.

				04 400 1	C 202 T	13.890	36, 308	7. 922	9, 590
	NT2RP3002659	18. 914	12. 170	24. 486	6.353			10, 430	27.532
	NT2RP3002660	64.465	53. 376	119.655	42.835	35.909	41.916		
	NT2RP3002663	30.048	20.813	29.457	13.786	13.557	15. 463	15.414	13. 216
5	NT2RP3002664	14.659	18.990	23. 494	8. 867	10.564	9. 625	5.085	4.798
	NY2RP3002667	15. 216	16. 234	11. 286	11.809	8.647	8. 484	26. 055	18.907
	NT2RP3002671	39. 495	26, 960	28. 177	15. 153	12.285	24. 589	13.809	14.654
		11. 347	14. 990	21, 206	28.999	14,002	9. 455	16, 128	33.677
	NT2RP3002682			16.858	8. 392	12.676	7, 181	5.777	4, 503
	NT2RP3002684	13. 722	11.697	10, 162	1.691	1.917	2. 141	3.706	3.397
	NY2RP3002687	2.560	4.651			13. 309	20. 413	8.939	1,088
10	NT2RP3002688	15.864	2.884	22, 879	1, 260			15. 073	59. 183
	NT2RP3002698	28. 485	12.350	29. 970	11.179	18.339	22.012		
	NT2RP3002701	144.580	68. 552	65. 738	22.713	47.971	117. 171	58.063	64. 453
	NT2RP3002705	50.811	34, 865	76.689	77.242	38.688	84.791	28. 441	54. 479
	NT2RP3002708	107, 193	25, 745	48, 335	10,739	20.147	29.081	22.130	32.554
	NT2RP3002711	38. 410	19, 460	31.129	24. 261	21.934	31.711	19. 413	9, 154
15		127. 597	337. 217	172.297	85. 410	157.291	209.750	71.600	90. 235
15	NT2RP3002712			26.653	9, 930	11.236	16.757	16.310	18.652
	NT2RP3002713	25. 722	12.997			19, 153	24. 353	19. 280	10, 413
	NT2RP3002721	48.039	15. 327	24. 924	23. 105		259. 199	308, 568	136.618
	NT2RP3002722	421.087	147.659	338. 772	115.647	164. 233		121. 373	45. 387
	NT2RP3002723	43.086	85.012	67.010	38. 528	35. 204	150. 941		16, 796
	MY2RP3002737	71.494	27.672	52, 178	22.716	32.049	58.862	47.802	
20	NT2RP3002738	47. 542	16.654	36.964	9.362	16. 223	38. 458	25. 360	23.198
20	NT2RP3002742	81.782	149. 322	102.776	54. 228	44. 909	105. 384	127.394	33.680
	NT2RP3002744	2. 263	4, 168	21.735	2.015	3.502	1.976	2. 225	0.560
	NT2RP3002756	22.619	12. 182	21.840	8.009	10.135	11, 380	12.917	4.838
	N12RP3D02757	113.772	65. 294	69, 951	34, 431	19.743	281.518	37.409	82.637
	NT2RP3002758	60.176	82.911	68. 360	23.774	51.197	81.519	55, 695	20.674
		70.007	62, 402	96,808	44.296	70.524	111.844	35.008	61.053
25	NY2RP3002762			93.384	42.890	27, 102	55.601	31.878	35. 587
	NT2RP3002763	65.632	38. 286			10.042	24.469	17.980	11.225
	NT2RP3002770	35. 381	13.511	35.913	7, 950		25.254	18. 920	17.572
	NY2RP3002771	40.863	23, 186	29,004	13.976	35.897		5, 889	5. 289
	NT2RP3002785	13. 960	5, 890	4, 173	2.677	2.677	9.071		
	NT2RP3002790	34.782	20.599	28.673	15.987	14.483	19. 288	18. 105	19.768
	NT2RP3002799	39,751	31.026	83.485	29.150	23.866	22.566	21.257	45.619
30	NT2RP3002801	47.659	26, 163	128, 555	31.073	22.498	25.337	24. 586	24, 190
	NT2RP3002802	146. 487	73, 131	121.221	33.065	38. 992	67.510	59. 237	21.826
	NT2RP3002810	10.160	45. 362	22.360	7.561	8.729	7.648	14. 315	7.654
	NT2RP3002818	4.667	6,464	10.095	3. 200	6.216	4.871	5.874	11.909
	NT2RP3002821	76.117	34, 802	53.630	32.950	28.735	58, 082	41.128	16.704
	NT2RP3002823	11, 784	13.818	14. 562	1.977	7.384	12.328	6.617	11.325
			13. 354	18. 512	8.300	12.766	20.235	20.838	24.852
35	NT2RP3002825	51.146	38. 250	97.142	25.989	24. 214	26.885	16.084	21, 503
	NT2RP3002829	35. 187		68.213	17.668	23. 336	61.962	46.206	37.479
	NY2RP3002831	66.496	27. 156			36, 995	100. 291	59.703	56.686
	NT2RP3002836	130, 172	72.920	90.667	20.404		22.445	10.215	12, 196
	NY2RP3002845	64. 337	22.726	40, 173	14.166	18. 291			7. 989
	NT2RP3002852	38.556	19.001	25, 493	7.993	8. 490	26.043	16.609 8.852	2.388
40	NT2RP3002861	2.544	8, 478	5, 538	2. 371	1.076	11.828		17, 114
70	NY2RP3002869	119.363	36.492	65.104	27.751	37. 200	48.198	61.052	
	NT2RP3002874	24.807	10.169	15. 126	5. 983	8. 446	11.486	15.977	17.599
	NY2RP3002876	64.967	22, 806	49.911	23.937	25.658	54. 137	50.714	12.582
	NT2RP3002877	86.753	69, 686	258. 276	48.444	44, 144	53.777	36.801	48.742
	NT2RP3002887	32,513	9, 192		15.590	7.085	25.821	19.262	5.065
	NY2RP3002900	17. 592	22.036			17.946	18.936	16.030	15. 494
45	NT2RP3002902	77.119	87.651				49, 857	35. 525	68.682
	NT2RP3002909	651, 498	271.044					375.523	192, 134
								14.454	11.143
	NT2RP3002911	18.365	31, 404	44.303				12.470	4.969
	NT2RP3002948	31.554	19, 471						9.777
	NY2RP3002953	86. 292	18.063						
	NT2RP3002955	19, 801	7.571						
50	NT2RP3002958	41, 536	22. 160	22.74	5.690				
	NT2RP3002969	37. 280	28.189			18.977			9, 514
	NT2RP3002972	22, 208	18.736						7. 568
			15.240						
	NT2RP3002978	17.816							
	NT2RP3002983	7.404	5. 940	7.10	2 1.188	1.140	3. 403	0.213	

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Table 106

NT2RP3002988 17.700 17.268 27.888 13.345 13.104 15.971 19	5. 968 9. 252	20.623
	9. 252	
		22. 620
	3. 309	47.982
5 NT2RP3003008 40.397 31.290 39.838 8.641 14.630 27.543 20	0.015	10.881
	6.941	4. 141
NT2RP3003015 54, 108 13.725 29.619 7.455 12.688 24.800 30	0.124	11. 125
	6. 155	2.761
	3. 397	10.834
	6. 530	32.637
	9. 952	17. 929
	1.309	0.000
	7.766	20.617
	5.007	75.424
	8.869	21.064
	2.042	87.952
7 20 10 10 10 10 10 10 10 10 10 10 10 10 10	5.012	5. 335
	3.431	12.628
	8. 951	10.008
	5. 965	42. 507
	4.319	136. 293
	7.416	13. 913
	9. 853	16.047
20	8.075	11.570
	7.673	8.566
	1.069	6. 398
	6.056	8.843
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	7. 334	18. 974
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M12A73003133 11.001 3.814 23.401 0.320 11.110 14.000 1	6. 054	10. 575
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	6.040	16.059
	29. 781	16.540
	0.660	81. 294
	32.807	25. 906
	24. 359	47.838
111 211 444 444 444 444 444 444 444 444	23. 116	10.543
	77. 380	29. 340
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IN CAN DECEMBER OF THE PROPERTY OF THE PROPERT	33.588	22.928
33	24.899	16.702
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N. 2.11. 000 00 00 00 00 00 00 00 00 00 00 00 0	6.674	9. 347
	13. 939	5.004
	14. 491	4. 525
	24.772	17.016
40 NTZRP3003242 16.643 9.743 12.011 3.953 5.705 9.943	7.847	1.554
	42, 132	48.708
	42.743	34. 203
	13.828	62.410
	31.572	22.039
	22. 285	16. 139
45 NT2RP3003273 11.164 9.672 10.474 15.421 5.945 12.757	7. 385	3.145
	10. 155	3.221
	21.479	28. 154
	27. 454	26.077
MT2AP3003290 149, 162 75, 603 249, 880 57, 514 56, 137 81, 416	57.703	30.573
	26.294	25.862
1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -	18.752	19.610
50 N12RP3003302 45.288 23.590 92.158 17.983 13.001 23.342 NT2RP3003311 4.124 7.411 10.651 5.453 14.885 11.665	3. 658	3.020
NT2RP3003312 14.814 8.617 14.507 5.774 2.403 16.774	9, 193	8.645
MT2RP3003313 15,411 6.290 9.374 4.661 3.186 10.303	5. 674	15. 392
	15. 226	21.848
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Table 107

	NT2RP300333D	29.506	12.597	10.896	8. 585	8.115	8. 559	6.939	9. 940
	WT2222244	29.694	14.023	28.467	10.446	14.551	23. 190	14, 110	21.136
	NT2RP3003344				37. 233	38. 412	50, 911	50, 114	35. 893
	NT2RP3003345	105.530	88. 425	241.668				15. 004	13.774
5	NT2RP3003349	20.318	21.037	19. 247	6.025	8. 572	15. 104		
-	NT2RP3003353	10.529	10.306	3.139	3.872	5. 195	16. 793	3. 277	2.796
			242.452	577.215	170.336	177.749	307.555	235. 179	214. 175
	NT2RP3003354	481.127				15. 329	29, 997	27. 654	13, 096
	NT2RP3003368	47.684	23. 833	38.838	12.045				11.087
	NT2RP3003375	9, 531	13. 959	20,610	8.653	7.770	15. 597	5.760	
		166.751	42.971	84.536	25.743	44. 033	73.870	73.821	25. 200
	NT2RP3003377			37.902	18.516	20.006	33,001	24. 969	18,065
10	NT2RP3003384	44.335	23.396				68. 473	67.072	48.712
	NY2RP3003385	94.843	42.782	74.715	20.456	31.187			
	NT2RP3003396	33, 482	30.352	33.756	14.143	15.615	30.475	16. 101	16. 251
			37.215	59.716	18.488	19.630	41.023	7.020	14. 203
	NT2RP3003403	53.313			10.044	13.315	26.899	23. 574	10.007
	NT2RP3003409	34.343	23.644	29. 939				32.903	26. 101
	NT2RP3003411	79.480	70.920	90.615	61.424	39.085	48. 593		
	NT2RP3003420	61,545	52.479	134.682	28.549	32.168	25, 103	23.751	18. 844
15			18.577	22.890	8,071	10. 241	21.558	25. 924	-11.363
	NT2RP3003425	28.870				32.319	90.461	44.692	26. 808
	NT2RP3003426	126.098	63. 120	93.804	24. 452				24.813
	NT2RP3003427	53, 936	61.645	67.284	18.467	14.098	40. 425	41. 425	
	NT2RP3003433	97, 022	87.577	196.547	46.930	103.713	35. 421	49. 581	51, 308
			90.341	101.893	38. 490	90.843	65. 265	43.848	39. 524
	NT2RP3003437	70.471				57.030	82.442	40.878	33.734
	NT2RP3003448	166.318	99.558	171.792	33. 106				
20	MT2RP3003455	98. 805	99.945	87.828	44.898	40.079	47,665	54.700	42. 051
	NT2RP3003462	42.184	21.903	23.018	11.812	14.369	18.994	22. 972	14. 965
			19.800	20.515	13.066	11.398	11, 185	9.509	8. 151
	NT2RP3003464	20.285				22.567	43.698	25.742	22.878
	NT2RP3003469	63.020	31.314	45. 443	12.277				60.344
	NT2RP3003473	49, 194	61.265	73. 244	52.029	33.239	49.762	41.082	
	MT2RP3003474	25. 607	8,816	7.783	3.674	4,629	13.456	6.864	6. 240
05			28.799	37, 252	11.016	19, 935	32.908	31.492	21.824
25	NT2RP3003475	68.962					3.738	7.208	8.419
	NT2RP3003490	20.464	20.731	22.026	3.717	16.041			9. 848
	NT2RP3003491	10.282	25.486	15.580	15, 193	6. 202	6.287	6.927	
	NT2RP3003493	225.729	58.149	69.338	48. 207	44.647	93.915	53. 796	47.878
			21.791	23.783	12.174	8, 905	10.384	6.189	9. 984
	NT2RP3003500	16.211				9.823	21.336	14.921	8, 623
	NT2RP3003527	35. 235	13.032	16. 125					14. 186
30	NT2RP3003532	35. 952	35.805	89. 452	21.080	32. 372	12, 131	23.670	
••	MT2RP3003535	30.511	17.215	16. 247	3.432	9.615	14, 199	11.449	7.658
			11.045	31.565		18. 265	21.717	21.923	38.703
	MT2RP3003536	35.415				40.066	19.654	56.835	72.031
	NT2RP3003543	69.871	52, 348	78. 481				31.229	23.611
	NT2RP3003549	42,025	14.802	50.570	18.842	33. 282	15, 787		
	HT2RP3003552	4.529	4.296	2.807	0.000	4, 647	10.319	2.766	9.014
			40.350	57.743		32.961	12.721	42.457	36.766
35	NT2RP3003555	57.410					5. 159	6.000	8.501
•	NT2RP3003559	20.086	11.398	15. 254				25, 239	22.138
	NT2RP3003564	66.462	28.214	41.863			36. 338		
	NT2RP3003572	50.882	28. 277	31.870	11.128		36, 904	28. 134	19. 912
	NT2RP3003576	236.584	162.700				90.587	262. 925	
							5. 529	28.863	23. 236
	NT2RP3003587	34.277	96.685				58.468		
	NT2RP3003589	69. 284	36. 270						
40	NT2RP3003592	93, 627	36. 255	60. 268			27.570		
	NT2RP3003593	64. 187	68. 925			11.913	10, 024	11.351	
									77.77
	NT2RP3003614	202, 651	80. 341						
	NT2RP3003621	15, 164							
	NT2RP3003625	131, 346	86.62	204.03	32.07				
	NT2RP3003627	95, 853				3 43.349	33.276		
45								40.566	25, 499
45	MT2RP3003636	87.887							
	NT2RP3D03642				3 (3.14)	27.04			
	NT 2RP 3003645	42.276	23. 45	37.01					
	NT2RP3003648				5 13.95	4 21.37			
									21.889
	NT2RP3003649								
	MT2RP3003650	70. 844	54.07	7 30.99					
50	NT2RP3003656			9 21.96	7 19.08	2 28.00			
30	NT2RP3003659						33.70	2 21.32	11.457
	NT2RP3003662								
	NT2RP3003664	31.48	1 40.03						
	NY2RP3003665				3.21	0 5.22	8.90	0 22.76	9 15.66
	W15V13003003								

Table 108

		•		140	16 100				
	NT2RP3003671	19, 991	15, 142	32.517	10.512	26.620	6.813	15. 367	4. 134
	NT2RP3003672	59.637	70.861	52.702	21, 219	42.465	28,220	33.602	25. 472
				29, 196	8. 319	9, 184	13.218	19, 475	8, 563
	NT2RP3003673	22. 381	26.615			55. 109	70. 199	47.217	161.678
5	NT2RP3003679	210.406	183. 454	88.575	68. 184			9.675	4. 489
	NT2RP3003680	36.432	9.725	11.980	2.868	17.580	10. 982		18.094
	NT2RP3003686	23.300	18. 187	60.813	12.758	15, 373	14. 321	13. 248	
	NT2RP3003689	16.292	10.228	7.344	18.943	22.892	23.049	1.755	14.648
	NT2RP3003697	18.041	18.889	23,041	11.465	5. 304	14.646	14.387	27.765
	NT2RP3003701	23.411	19. 362	26.737	5. 128	15, 953	19.109	21.561	12. 363
	HT2RP3003704	83.293	69.818	227.532	48.512	34.531	23.793	34.747	31.728
10	N12RF3003104	27. 845	26.701	35.031	16.071	7.707	11.396	10.568	7. 157
	NT2RP3003714			32.116	1, 957	10.013	19. 271	16.236	6.862
	NT2RP3003716	23. 382	29.412		16. 267	34.684	31.096	35.668	28.013
	NT2RP3003721	47, 677	30. 191	49. 197		14. 255	9. 224	6.260	18, 801
	NT2RP3003722	23.636	24.625	30, 510	14.691			45.483	33. 548
	NT2RP3003726	71.518	25. 344	63. 123	17.350	34.451	43. 109	29, 613	40. 648
15	NT2RP3003729	48.252	22.558	41.664	11.182	23.933	14.940		
	NT2RP3003731	117, 126	53.921	150,601	44. 104	59.737	67.883	52.971	70.102
	NT2RP3003740	95. 127	38.608	55, 360	23.461	31.988	57. 694	54. 566	25. 167
	NT2RP3003746	16.191	12.220,	16.980	7.510	13.596	12.664	8.103	4.718
	NT2RP3003749	0.000	0.000	0.000	0.603	0.000	2.487	2.557	3.945
		15.865	21, 394	19, 162	12.449	13.299	26.475	9.854	18.648
	NT2RP3003754		0.000	0.000	1.040	0. 228	0.000	0.000	0.000
20	NT2RP3003759	0.000		64.694	. 34. 845	35. 239	58. 222	58.654	59. 695
	NT2RP3003764	83.938	66.804		12.627	24.046	19.839	39, 865	29, 001
	NT2RP3003766	65.630	30.349	55. 241		34.723	31.155	25.595	43.641
	NT2RP3003767	70.910	69.657	250.723	42.998			54. 893	62.981
	NT2RP3003778	131.825	86.793	385.771	86.755	57.514	68.379		45. 498
	NT2RP3003779	109.510	79.471	82.764	30. 193	42.973	68.003	45. 497	
	NT2RP3003783	20.728	49.548	65.851	31.076	42. 337	19.891	30, 990	36. 918
25	NY2RP3003787	52.420	24.376	34.398	5.999	3.586	110.807	52.440	37.987
	NT2RP3003789	49.434	35, 220	51.425	19.152	23.911	36, 130	35. 358	51.169
	NT2RP3003795	35, 141	27.549	49, 460	9.850	9.646	24.082	23. 805	22.055
	NT28P3003799	43, 365	13.905	22.874	6.981	14.894	24.044	24.707	15. 462
	NT2RP3003800	33.918	17. 363	27.230	9. 216	12.645	25. 354	23.431	31.197
		63, 293	44.084	37, 398	25.212	22.134	20.827	35.180	33.836
30	MT2RP3003805		50. 351	23.357	8.497	6,068	18.501	12.588	23. 510
30	NT2RP3003809	31.815		386. 972	66.656	124,750	204. 320	163.951	105.623
	NT2RP3003819	524, 121	195. 245		9, 543	22.963	19.518	18, 840	18. 478
	NT2RP3003824	23. 645	17.797	34.795		51. 197	72.544	45.338	78.057
	NT2RP3003825	100. 544	64. 212	102.915			6. 483	4.696	6.839
	NY2RP3003828	13.857	3.284	8, 953	5.968	12. 172		36.027	61.956
	NT2RP3003831	58.812	63.105	141.638		42.372	35. 689		29. 782
35	NT2RP3003833	37. 263	25.079	32.114		15. 132	21.745	17. 267	
	NT2RP3003836	139.979	72.806	102.049		60.838	71.273	62.037	67.712
	NT2RP3003842	173.727	172.520	421.266		82.994	67.844	51.328	70.400
	NT2RP3003843	40.446	57.570	27.866		61.585	12.265	18.777	39. 377
	NT2RP3003844	71.843	59, 271	53. 342	25.835	23.638	29.874	45,658	29, 555
	NT2RP3003846	9.016	12.338	29, 501	8.508	8.017	9. 155	11,844	13.878
	NT2RP3003849	59. 374	29.253			18.400	31.563	24. 824	35. 683
40	NT2RP3003862	28.859	32.198			14.207	16.311	10,540	19, 157
	NT2RP3003870	163.978	56.534				66.418	66.181	41.207
	NT2RP3003874	25, 106	64. 501	32.262			22.879	79.189	8.302
							19, 085	11.236	26.223
	NT2RP3003876	57.365	29.873				26.939	26,220	22.845
	NT2RP3003880		23. 356				9. 987	0.000	44. 372
45	MT2RP3003889	7.749						10.999	21,695
45	NT2RP3003891	25.663							
	MT2RP3003914	84.860					38.079	39.405	
	NT2RP3003915	24.657	11.712					22.731	
	NY2RP3003918		28. 378					27.234	
	NT2RP3003920			182.18					
	NT2RP3003924					20.859			
50	MT28P3003932						15. 130		
	NT2RP3003939							16.204	23.960
									32. 163
	NT2RP3003940								
	NT2RP3003943								
	NT2RP3003959	33.09	7 24.51	8 31.71	3 1 13.33	13.31			

Table 109

					I	(3.060.1	90 205 1	01 146	44.991
	NT2RP3003963	225. 975	65. 265	81.733	29.808	52.069	80. 205	81.146	123. 379
	NT2RP3003965		148.769		23.378	65.718	64.058	36. 726	52.120
	NT2RP3003972	178.647	135.585	147.168	34.841	77.695	106.673	70.941	45, 075
5	NT2RP3003973	62.806	37.262	47.172	25. 442	23.541	30.764	27.857	
	NT2RP3003979	42. 205	32. 192	109.653	39.966	32.734	35, 850	18. 262	64. 857
	NT2RP3003980	43.589	24. 631	25.030	11.906	6. 253	21,641	13. 122	23.086
	NT2RP3003982	12.297	22.386	11.608	2.387	11.030	5.747	12.456	34. 995
	NT2RP3003989	17. 308	4.219	22.495	7.718	11. 234	3.600	3. 546	106.880
	NT2RP3003992	38. 217	23.384	39. 566	7.169	21.356	24. 091	21. 385	25, 954
10	NT2RP3004000	14.260	12.046	9,623	3, 141	15. 292	10.563	26. 334	5. 687
10	NT2RP3004001	15. 524	17.005	53.914	11.406	10.314	27. 264	13. 462	16.712
	NT2RP3004005	9.869	9. 263	84.786	19.372	0.000	4. 857	1.497	9.756
	NT2RP3004013	14. 485	12.461	42.406	11.492	13.049	8.125	6.478	17.758
	NT2RP3004016	26. 353	20.174	14. 242	8.659	7.098	11.464	20. 928_	17. 553
	MT2RP3004025	60.555	22.329	39, 729	22.559	18.276	23, 525	24. 555	35.771
	NT 2RP 3004023	612.399	230. 471	834. 283	175.098	230. 371	417.549	400.971	300. 584
15		35. 758	19.204	29.889	17.016	20.612	23.674	15.019	17.667
	NT2RP3004041		150. 283	197. 509	53.931	78.902	154, 218	126.411	98.212
	NT2RP3004042	212.341		51.482	17. 239	26. 945	24.143	30. 198	21.882
	NT2RP3004044	72.252	110.791	142. 232	35. 932	51.071	43.163	38.869	49.345
	NT2RP3004051	152.863	73.839	74.633	29.148	35. 481	72, 900	21.817	40.892
	NY2RP3004052	121.021	59, 192	277.692	61.036	78.666	68.730	38. 992	75.069
20	NT2RP3004053	98.068	91.523	20.623	13.216	5. 886	21,414	72.807	7. 926
	NT2RP3004055	94.456	63.815		33.539	23.030	10.773	12. 908	18.849
	NT2RP3004059	26.860	40.017	21.750	7.097	24. 752	5, 609	24.116	30, 966
	NT2RP3004063	18. 643	7.895	20. 299	48.083	19.941	65, 794	83.498	20.778
	NT2RP3004067	252. 237	73.282	95. 895	21, 957	33.015	28. 191	23.513	30. 233
	NT2RP3004070	48. 573	60.633	86. 573	11.710	25. 118	31.470	27.043	31.641
25	NT2RP3004075	38.601	29.096	32.376	18.027	27. 424	76.975	68, 265	35.076
25	NT2RP3004078	123. 241	42.946	72.005	10.656	16. 243	25. 486	10, 927	25.077
	NT2RP3004083	44. 275	15.592	19. 299	13.491	18. 285	6. 670	5.617	3.170
	NT2RP3004084	20.841	11.260	17.316	34. 544	41. 231	18, 188	46, 470	43, 578
	NY2RP3004087	61.884	66.963	88. 119	21.173	17. 529	18.879	17.880	26. 579
	NT2RP3004090	36.365	32.568	40. 579	50. 577	97. 795	88.393	53, 404	59. 593
	NT2RP3004093	161.528	139.905	344. 325	60.637	74.060	107.607	74. 457	93, 441
30	NT2RP3004095	200.143	125. 167	292. 455	25.857	52.758	90.150	84.250	44,710
	NT2RP3004102	189.415	73.338	84. 114		74. 491	121. 974	73.119	123. 538
	NT2RP3004110	147.625	133.897	357.078	89.105	44. 915	38.873	58. 991	47. 932
•	NT2RP3004119	104.164	75. 262	197.706	41.775	117. 997	203. 963	194. 543	177. 494
	NT2RP3004125	312.772	144.655	288. 945		13.862	6. 399	91.521	13.988
	NT2RP3004129	32.046	25. 525				35. 035	32.730	28. 345
35	MT2RP3004130	49. 467	45.820			28.933	33. 261	34, 485	27. 866
33	MT2RP3004133	55. 970	58.961	100. 212		9. 248	49, 189	43.159	26. 374
	NT2RP3004145	105.806	51.341	52, 276		30.673 47.130	133.546	97.568	36. 471
	NT2RP3004148	206.658	51.505				47. 245	42.482	35. 181
	NT2RP3004155	65. 340	68.555			55.725	39, 412		33, 175
	NY2RP3004165	31.599	44.217			20. 207	24, 234		
	NT2RP3004179	35. 856	20.632				12.464		14. 309
40	NT2RP3004185	32, 929	15.710				61, 530		
	NY2RP3004188	125.817	53. 211			32.369	45, 470		
	NT2RP3004189	71.207	30.246				16, 769		
	NT2RP3004190	23, 559							
	NT2RP3004191	83. 281					25. 270		
	NT2RP3004202	65.428							
45	NT2RP3004205	85.092							
	NT 2RP3004206	14.256							
	NT2RP3004207								
	NT2RP3004209		24.20	3 39.56					
	NT2RP3004215				9 8.18				
	NT2RP3004219					51.38			
50	NT2RP3004242						15.11		
50	NY28P3004246							32.83	
	NT2RP3004253							0 29.63	2 32.520
	NT2RP3004258								
	NT2RP3004262	71.43	4 29.97	41.00	12.02	4 4 4 9 1		1	

Table 110 ··

		00 F00 T	36.290	83.006	24.540	22.746	61.823	54.050	37.950
	NT2RP3004275	98. 599		178.061	49.657	96.836		106. 109	47.357
					7. 292	8.756	13.882	7.956	36. 428
	NT2RP3004289	15.745	32. 192	24. 193		5. 484	19.565	13.721	12, 117
5	NT2RP3004294	60.266	26.724	26. 421	11.149		81, 222	91.861	86, 967
	NT2RP3004298	132.592	61.132	108.061	41.028	51.835			51. 104
	NT2RP3004309	144.028	38.007	72.661	18.449	49.804	89.984	72. 157	
	NT2RP3004321	231.684	53.180	108.237	29.746	51.266		104. 335	90.745
	NY2RP30D4322	37.875	23. 343	25.724	12.249	19.668	22.470	23.599	36.486
	NY2RP3004332	106.333	91.471	249.231	44.955	55. 341	76.389		107.059
		68.850	32.416	38.730	9.752	18, 775	14.058	18.048	16.320
10	NT2RP3004334		59.827	77.110	20.736	37.630	26.664	34. 386	34. 983
	NT2RP3004336	51. 294	16.241	17.569	3.872	10.946	14.386	14, 110	86.362
	NT2RP3004338	18. 622		19.614	6.657	8. 502	12.520	6. 268	32.744
	NT2RP3004341	19. 200	20.230		9.060	9.720	11.640	14.563	16.985
	NT2RP3004345	23. 525	19.497	30. 403		108.792	59. 212	48.175	79.425
	NT2RP3004348	152.635	117.901	359.204	67.822		53.467	43.103	73.727
15	NT2RP3004349	156.222	104.964	468.032	69.388	77.765		127. 957	121.148
13	NT2RP3004355	58.395	30.712	72.395	19.596	16. 476	48.617		48.103
	NT2RP3004356	110.831	61.735	75.603	20, 147	52.762	88.239	65. 266	
	NT2RP3004360	41.674	35. 467,	41.306	19.910	12.453	15.566	22.989	27.326
	M12RP3004361	46.996	33.404	30.049	14.201	14. 577	23.509	11.195	14.723
	HT2RP3004374	95. 389	57.120	48. 566	15.283	39. 161	43.002	46.264	23.628
	NT2RP3004378	58.764	49.662	50, 107	18.157	38. 127	30.939	38.526	49.715
20		16.800	27. 122	23.992	18.781	27.931	12.441	19.782	23. 253
	NT2RP3004399	76.975	42.401	68.536	14, 461	40. 127	30.855	27.361	25.603
	NT2RP3004405		18.451	36.531	9.936	27.693	43.690	31.470	25. 327
	NT2RP3004406	59. 371		74, 904	12, 415	33. 625	61.907	28.318	22.563
	NT2RP3004411	92.442	48. 901	29. 952	10.559	13.320	23.158	18.753	13.677
	NT2RP3004424	40.886	26.604		18.251	39.655	61.213	57.747	33.647
	NT2RP3004428	141.707	50.415	59. 329		175.041	22.755	14.727	14,260
25	NT2RP3004432	26.049	27. 127	235.751	18.465 32.310	42.640	67.791	64. 267	46.448
	NT2RP3004434	146.690	70.435	71.916			5. 257	19.506	10, 316
	NT2RP3004446	27.192	19.189	44.272	8.673	16. 147	15. 972	19.748	17, 124
	NT2RP3004451	45.826	25.986	81.355	14.858	17. 991		8.872	9.047
	NT2RP3004454	13.596	21.506	24, 434	5.907	6.024	8.062	173.844	118.891
	NT2RP3004466	267.157	127. 933	175.917	65.272	67.867	153. 148		95, 231
30	NT2RP3004470	150. 361	134.643	271.527	54.812	70.601	50.612	49.084	25. 525
30	NT2RP3004472	13.995	10.444	6.945	8.463	7.742	9, 150	3.258	
	NT2RP3004475	89.313	39.845	56.364	22.197	34.071	46. 397	52.228	36.349
	NT2RP3004480	27.508	23.946	28.297	14.978	36.756	18.216	23.949	28.732
	NT2RP3004481	31.506	22.386	32.532	15.846	17. 215	13.188	11.393	75.655
	NT2RP3004490	5. 922	2.592	0.000	0.000	0.000	0.000	8, 285	6. 521
	NT2RP3004496	24.027	28.908	28.749		13.349	15.561	11.595	12. 252
35		109.432	51.964			34.097	43.928	34.988	37. 439
	NT2RP3004498 NT2RP3004503	162.798	115.770			66.406	56.670	46.593	74,722
		62.371	28.837		18.389	15.784	30.245	70.081	29. 325
	NT2RP3004504		46. 920			11.983	15.997	28.823	36.454
	NT2RP3004505	25, 650	32. 594			25.414		34.107	31.896
	NT2RP3004507	50.531	14. 576			25. 541		6.345	25. 622
40	NT2RP3004519	38.355	27.009					33.495	24.924
40	NT2RP3004524	38. 228							15.761
	NY2RP3004527	27.651	20. 933						9.045
	NT2RP3004534	33.516	8.840						73.442
	NT2RP3004539	100. 285							12.869
	NT2RP3004541	35.828							94. 341
	NT2AP3004544	52.885							14.870
45	NT2RP3004551								
	NT2RP3004552	100.028					1 1 10		
	NT2RP3004557								
	MT2RP3004561	103.770	34. 28						
	NT2RP300456								
	NT2RP300456	94.551			3 22.78				
	NT2RP300457	55.49			9 14.634				
50	NT2RP300457			77.0		9 20.59			
	NT2RP300458					8 32.01			
	NT2RP300458								
							5 12.91	3 32.94	25. 495
	NT2RP300459	4   40.17							

520

Table 111

				24 323 1	47.277	28, 549	51.397	38.270	98. 212
	HT2RP3004603	78.679	80.544	62.737					21,608
	NT2RP3004612	74.014	32.975	30.756	11.218	37.649	29. 374	13.820	
	NT2RP3004617	34, 514	16.958	15.437	7.541	9.813	10, 352	13.498	6. 437
5	NT2RP3004618	45.654	67.084	24.650	10.899	12.856	27.696	15.781	34, 862
-	NT2RP3004625	75.276	30.663	96.644	20.740	43.066	82. 423	59. 145	28. 085
		67.742	53.096	56.701	30, 583	29.960	46. 122	44.888	61.643
	NT2RP3004635		58. 380	202.476	49.309	45, 610	45. 215	57, 393	54, 691
	NT2RP3004640	89.717		118.760	36.694	65. 566	113. 287	75.702	49.519
	NT2RP3004642	173.246	73.060			50. 982	53, 766	48.670	44. 858
	NT2RP3004647	101.143	79.944	113, 136	52.874		63. 735	70.579	53. 558
10	HT2RP3004652	203.591	158. 366	434, 477	72.065	120.412			
	N12RP3004669	83.602	70.489	66. 421	12.848	23. 192	58. 448	88. 231	37. 292
	NT2RP3004670	193.547	128.951	178.554	73. 935	102.781	166. 902	107.905	94, 007
	NY2RP4000008	19.767	47.505	24, 109	17.304	29. 354	55. 419	33. 855	34. 432
	NT2RP4000018	56.348	39.769	80.074	15.072	26.721	42.484	38.619	43.517
	NT2RP4000023	53.022	17.753	34, 758	10.911	23. 301	26.391	19.092	19.833
			56.593	72.466	8.582	83.053	47. 152	45. 373	52, 951
15	NT2RP4000025	45. 646		321, 911	40.713	60.319	94. 350	45. 943	45. 399
	NT2RP4000035	119.584	72.523			47. 224	60. 208	34. 302	31.401
	NT2RP4000041	186.503	56.255	41.691	8.801			24. 059	13.748
	NT2RP4000049	47.651	27. 923,	39.552	7.903	6.803	18.769		
	NT2RP4000050	46.861	18. 274	33.191	8.103	13. 428	12.029	13.779	7. 279
	NT2RP4000051	40.843	29.142	32.303	10.190	21.384	40.455	39.037	17.835
	NT2RP4000063	43.284	30.034	25.813	11.605	18. 431	28. 262	27. 310	20.178
20	NT2RP4000065	11.102	17.154	21, 158	43.890	19, 264	6.730	6.069	32.776
	NT2RP4000070	59.796	43.567	133.907	34.788	23.019	47.653	20.318	14. 552
	NY2RP4000074	18. 725	4.052	10, 370	1.424	4. 150	8.454	6.795	2.366
		62.113	86. 532	57.818	34.813	30. 151	56. 743	50. 257	36.799
	NT2RP4000078		111.931	192.627	75, 992	91.873	205.033	130.550	126.661
	NT2RP4000080	224.722		1600. 483	150.687	285.007	248.048	126.052	293.699
	NT2RP4000099	321.974	219.279				9. 573	53, 928	13.964
25	NT2RP4000102	8.753	18.572	15.774	4. 228	7.806		47.841	72.767
	NT2RP4000103	34.791	23.847	32.776	10.952	8. 411	17. 791		
	NT2RP4000108	62.537	43.717	44.931	25.841	148. 533	28. 159	30.906	35. 415
	NY2RP4000109	261.144	124. 505	231.410	69.135	84.528	232. 287	157. 290	146.451
	NT2RP4000111	28.240	10.956	13. 276	3.790	9.951	18. 128	12.668	12.698
	NT2RP4000112	174.823	126.761	222.355	29.525	41.360	94. 077	68.016	67.817
22	NT2RP4000115	104.464	46.026	87.051	17.566	38. 187	78.479	43. 365	44. 515
30	NT2RP4000129	20. 582	20.434	22.054	7.476	11.813	11.733	11.125	12.513
	NT2RP4000137	40. 931	26. 333	38.192	19.805	13.933	28.819	22.933	25.032
			41.054	56.796	8. 100	30.556	62. 995	15.210	44. 386
	NT2RP4000138	53, 828		27.517	15.337	27.602	16.576	20.734	34.135
	NT2RP4000141	62.206	42.856	24.754	8. 363	10.418	21.963	32.513	27.229
	NT2RP4000147	26.467	16. 245		111.407	84. 297	120. 085	78.831	153, 213
35	NT2RP4000150	170.729	155. 621	193.591		34. 976	55. 423	46.381	38. 147
00	NT2RP4000151	89.499	70.326	88.485	15.693			142.467	214. 943
	NT2RP4000157	374.212	306.778	1320.234	101.052	267. 293	258.633		
	NT2RP4000159	21.294	38.510	22.222	4.978	9.029	6.726	11.020	2.839
	NT2RP4000163	38, 106	28. 442	47. 497	14. 252	14.961	40.800	33.454	23. 270
	NT2RP4000167	20.173	26.500	23.216	7.845	5. 552	5. 423	7.245	14. 035
	NT2RP4000171	81.073	52.022	67.728	21.187	28.509	44.872	35.093	37. 752
40	NT2RP4000175	81.743	84.274	82,433	36, 175	79.980	58. 585	86.742	88.656
	NT2RP4000180	58, 476	59. 435	73.494	30, 105	37.648	47.113	80.700	76. 984
	NT2RP4000185	92.601	101.645	150.266	44.577	77.183	75.717	60. 488	85.600
	NT2RP4000192	127.476	49. 521	75, 782		46.143	55. 129	61.367	32.097
		56. 167	54. 180	31.757		23.917	32.670	26. 241	35.726
	NT2RP4000194						42.066	78.755	73.674
	NT2RP4000196	92.478	57. 125				369. 938	361.357	310,071
45	NT2RP4000210		304.062					87.129	138.067
	NT2RP4000212						119.552		
	NT2RP4000214	209.094	145. 483	438.818			69. 191	73. 163	99.829
	NT2RP4000216	27.754	23.804	32.743	9.142				26.648
	NT2RP4000218				25. 931	25. 141	34.742		62.428
	NT28P4000223							184. 145	106.046
50	NT2RP4000243								
	NT2RP4000246								
	NT2RP4000250								
	NT2RP4000256	61.500	54. 53						
	NT2RP4000257	146.739		68.08	1 16.98	74.826	29. 177	32.953	39. 299

Table 112

	NT2RP4000259	36.679	60.559	46.332	10.684	19.988	21.634	16.480	15. 511
	NT2RP4000261	43, 317	19.258	30, 162	7.462	9.311	20.800	15.617	17.669
				41.516	10.478	21.699	32.040	20.770	27.384
5	NT2RP4000262	57.147	28.869						
•	NT2RP4000263	26.287	13.027	49.010	13.046	27. 187	12.910	17. 489	13.293
	NT2RP4000280	404.385	153.579-	276.968	132.346	126.840	273.688	195.012	134. 292
			68.061	124. 456	10.943	103.023	163.664	158, 229	165. 646
	NT2RP4000286	349.970							
	NT2RP4000290	69.776	37.297	56.790	14.548	26.462	24.909	28.704	27. 597
	NT2RP4000291	92, 235	210.055	87.276	110.666	29. 297	73.542	109.583	151.177
	NT2RP4000301	72.312	25.823	43. 205	17.404	22.667	20.721	34, 359	47.720
10									
	NT2RP4000312	30.600	23.813	38, 345	71.709	0.000	27. 976	30. 543	16.077
	NT2RP4000321	152.139	101.314	320.889	47.164	45.419	56.735	18.656	58. 799
	NT2RP4000323	37.462	25.699	95.138	15.085	31, 924	10.455	5.460	17. 376
						54.837	40.659	43. 151	23. 155
	NT2RP4000324	336.502	41.027	28.832	17.302				
	NT2RP4000334	115.354	138.505	182.550	93.928	63.038	90.617	72.433	115.991
	NT2RP4000343	75.003	25.817	17,727	13.013	26.022	34.661	24.607	19.361
15	NT2RP4000348	56.032	12.454	12.331	15. 203	15.484	6. 180	3, 506	18, 445
	NT2RP4000349	7.762	0.000	0.000	3.720	0.000	0.000	0.000	6.473
	NT2RP4000355	87.546	71.121	115.193	27.548	24.554	33.248	29. 345	30.833
	NT2RP4000356	211.845	121.033	114.259	51.743	65.136	144. 965	93.350	89.148
					10.374	34, 417	19, 318	20.576	39.379
	NT2RP4000360	70.699	38. 241	86.142					
00	NT2RP4000367	18,288	5.279	7.668	4.052	7. 149	4. 373	5.067	3. 767
20	NT2RP4000370	32.692	19.934	38.747	6.510	17. 936	9, 489	6.000	24. 412
		8.950	23.267	11.530	6.424	4. 499	3.890	0.839	4.844
	NT2RP4000373								
	NT2RP4000376	35.864	18.265	19.621	12.884	15. 395	5, 826	23.805	21.083
	NT2RP4000381	46.926	33.825	103.826	18, 455	27.076	17, 117	10.557	22. 372
	NY2RP4000388	5084.865	1317.306	2099.929	227.725	2132.319	3323.080	4907.667	1152.125
						85. 442	187.036	159.581	
25	NT2RP4000390	257.545	160.161	219.816	71.826				156. 149
23	NY2RP4000393	12.640	11, 957	20.415	9, 221	11.409	7. 438	11.324	8.524
	NT2RP4000398	17.518	22.875	62.033	33.290	29.094	38.274	16.243	64.756
		72.166	37.198	50.776	14.912	16.850	25.605	52.793	18.016
	NT2RP4000406								
	NT2RP4000407	17. 281	27. 203	36.363	15.988	14. 182	13, 109	11.945	14.661
	NT2RP4000413	28.139	4.608	24.755	4.471	18. 199	9.618	9.564	3.410
	NT2RP4000415	52.988	28. 236	62.216	11.670	19. 273	18.078	30.417	40.803
30					20, 336	52. 584	49.364	45. 494	40, 422
	NT2RP4000417	120,835	54. 541	46.666					
	NT2RP4000423	45.442	44. 179	39. 359	11.506	22.404	15.869	30.636	33.860
	NT2RP4000424	69.125	46.323	210.620	28.361	37.650	36.808	15. 234	39.788
	NT2RP4000447	43.171	50.572	84. 440	39.944	38.491	45.721	39.832	64. 904
	NT2RP4000448	19.367	24. 180	80. 917	16, 101	11. 296	3.059	13.254	21.512
	NT2RP4000449	13.620	10.795	11. 538	2.925	6.616	4. 388	8.988	2.997
35	NT2RP4000453	16.784	23.231	20. 252	12.639	17.714	8. 345	19.980	15.034
	NT2RP4000455	24. 141	9.211	25. 236	8.774	21.609	10.059	20.357	12.379
	NT2RP4000456	119.272	61.157	163.661	22.286	65. 150	132.301	52.249	54.831
	NT2RP4000457	64, 205	43.798	49.492	18, 495	31.270	76.065	78.938	18,719
	NT2RP4000461	24.023	16.736	42.860	8.086	28.640	24, 287	12.689	10.443
	NT2RP4000462	61.975	32.022	55. 648	25.804	23. 165	20.388	41.481	46.650
40									
40	NT2RP4000463	44.030	41.396	65.217	27. 109	26. 324	27.922	36.605	49. 391
	NT2RP4000471	37.502	19.098	33. 476	5.338	11.489	19.044	0.000	11.363
	NT2RP4000472	13.349	14.082	11.918	3.395	5.066	10.401	8.705	6.892
	NT2RP4000476	8.321	93.773	34, 435	13.728	23.669	4. 372	15, 350	7.001
	NT2RP4000480	211.458	95.964	129. 427	15.810	72.857	76.584	80.179	54. 430
	NT2RP4000481	31.888	26.600	25.630	7.943	9. 597	13. 290	14. 597	17. 385
45	NT2RP4000483	21,998	15. 487	14.048	11.756	10, 365	13.738	23, 308	15, 114
	NT2RP4000487	60.364		22.474	11.302	12.610	14.044	7.594	9. 748
			31.407						
	NT2RP4000495	5.856	1.759	0.000	1.020	0.000	1. 332	1,331	1. 300
	MT2RP4000497	14, 222	23.785	35. 435	9, 191	6.838	6.266	19.870	19.909
	NT2RP4000498	10.973	30.501	18.513	11.562	11.061	3.896	18.332	11. 258
						20.760	15.985	16. 224	
	NT2RP4000500	28.356	22.346	29. 213	6.186				7.833
50	NT2RP4000507	65.764	65.249	44. 910	11.415	12.964	62.638	27.083	16.799
	NT2RP4000515	326.302	155. 582	205.890	76.678	101.826	196.853	160, 500	152.025
		44.510		143.747	33. 380	28.078	31.697	20.743	51.511
	NT2RP4000516		41.687						
	MT2RP4000517	43.875	14. 219	143. 214	16.861	8. 127	16.458	9.150	20. 642
	NT2RP4000518	26.023	21.987	59. 276	7.160	16.049	11.817	12.546	27. 280
	NT2RP4000519	26, 153	8.810	13.853	3.109	6.990	8.139	7.151	18.564
EE				0.000	0.000	0.000	0.000	0.000	11,634
55	NT2RP4000524	1.938	0.000	1 0.000	1 0.000	0.000	1 0.000	1 0.000	1 11.034

Table 113

	1248664866598 T	12.526	60. 186	18.819	3.919	15. 244	19.800	6.732	22.213
	NT2RP4000528				45.816	89. 192	83.433	71.078	86.062
	NT2RP4000537		47, 194		11.695	21.855	33, 231	40.279	26. 263
5	NT2RP4000541	106.565			15.964	35. 981	45.932	36, 402	28.580
	NT2RP4000543	121.504	31.320		83.348	51.684	53.797	34. 347	94.961
	NT2RP4000545	109.666			21.240	28.754	7.806	12, 598	34.617
	NT2RP4000546	34.736	*****			24. 334	25. 452	35. 475	51.804
	NT2RP4000549	27.942	60.396	16.907	8.050	12.069	10.840	14, 194	24.088
	NT2RP4000556	22.418	10.709	22.462	7.923			15, 418	23.701
10	NY2RP4000557	22.285	18.841	21. 106	3.617	11.430	13.950	52.601	55.628
10	NT2RP4000558	98.220		112.943	14.814	42.417	80.107	88. 953	88. 195
	NT2RP4000560	145.648	126.576	198.616	29.117	67.842	111. 268		1,728
	NT2RP4000568	4.653	7.710	9. 495	4.212	14.707	5.118	4. 418 57. 364	56. 537
	NT2RP4000583	100.314	94.610	258.628	54.914	59.898	38. 219	12.368	9. 441
	NT2RP4000585	36.734	19.742	25. 585	3.609	10.851	9, 594		9. 648
45	NT2RP4000588	24.965	28. 422	24.615	3.894	8.655	9. 562	10.506	28.565
15	NT2RP4000590	82.643	29. 520	74. 380	7.381	16.388	16. 999	38. 929	
	NT2RP4000599	5, 134	12.959	2.254	2. 300	0.000	5. 232	2.076	4. 437
	NT2RP4000603	48, 331	23.244	35.033	10.422	23.763	77. 588	27.888	18.472
	NT2RP4000607	43.033	46.964,	51.845	3.610	170.311	14. 213	16.592	35. 286
	NT2RP4000614	93.469	104.724	288.948	65.946	55. 948	39. 332	42.871	69.619
	NT2RP4000634	41.268	55, 106	42.356	20.080	29. 301	16.909	25.716	34. 506
20	NT2RP4000638	38.714	37.491	60.350	10. 197	20. 301	7.339	21.773	11.532
	NT2RP4000648	28.051	19.136	29.021	11.429	52.517	8. 564	11. 255	17.817
	NT2RP4000657	59.641	34.960	39.531	15.723	16.922	9.859	13.485	21.954
	NT2RP4000691	25.254	56.069	53.527	20.960	17.701	25. 333	15.651	24. 709
	NT2RP4000697	41.565	23.570	47.024	8.681	17.064	41, 529	26.741	15.415
	NT2RP4000704	150. 527	58.692	94.083	27. 108	61.336	83.179	82.422	52.001
25	NT2RP4000710	544.068	385. 881	401.163	199.745	108.821	570.526	370.976	288,408
	NT2RP4000713	28.318	29, 133	25,800	8.247	17.041	12.819	13. 220	15.778
	NT2RP4000724	15.864	37, 851	33.515	4.863	0.000	12.161	11.700	21.516
	NT2RP4000725	73.250	28.340	42, 587	10.791	15.656	23.049	29.695	16.914
	NT2RP4000728	398. 420	264.734	679.544	140.230	76.304	191,521	224. 945	194.628
•	NT2RP4000737	10.955	3.270	11.232	3.668	5. 117	2.568	5.042	3.465
30	NT2RP4000739	15.887	23.255	23,005	9.500	14.336	12.603	11.904	11.565
	NT2RP4000749	66. 966	32.925	44.669	15.449	15. 178	33.005	27.405	18. 522
	NT2RP4000769	65. 261	48.013	75.648	22.094	24. 165	36.022	30.919	26.509
	MT2RP4000774	42.939	36. 592	46.497	13.414	18. 307	19. 211	16.686	12.228
	NT2RP4000781	34.651	17.546	33.740	8.360	9.849	17.872	14.911	6. 625
	NT2RP4000783	29, 279	12.391	20.881	15. 327	3.867	20.509	21.416	4. 930
35	NT2RP4000787	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
	NY2RP4000788	57. 142	47. 566	42.475	22.374	15.545	36.822	18.884	32. 902
	NT2RP4000792	26.349	10.430	22.784	9.272	0.000	13.445	11.068	16. 223
	NT2RP4000809	33.934	109.004	47.604	14.815	14.118	130. 537	459.568	2.963
	NT2RP4000817	76.682	20. 256	38.151	11.596	23.415	26.562	17.001	12.542
	NT2RP4000821	121, 213	96.900	50.576	24.242	27.444	74.033	37.727	20.369
40	NY2RP4000822	140.413	82. 390	238.604	35.669	42.569	28.697	55.099	10.656
	NY2RP4000823		158.604	92.017	60.055	51.992	105. 428	517.857	15.029
	NT2RP4000831		29. 385	59.567	15, 141	27.742	44.635	56.751	39.831
	NT2RP4000833		143. 283	293.871	54. 134		59.985	36.700	53. 259
	NT2RP4000837		62, 893	85.421	24.336		108.156	63.147	18.389
	NT2RP4000839		59.635	88.717	37.592		49.470	22. 530	10.946
45	NT2RP4000846		31.507	17.224	17.876		20. 275	20.877	18.662
.•	NT2RP4000848		69, 956	299.625	42.214		34, 689	23.120	18.618
	NT2RP4000855		17.013	12.507	9. 287	13.091	8.594		11.970
	NT2RP4000863					3. 763			4. 230
	NT2RP400086								
	NT2RP400087					40, 443			
50	NT2RP4000874	114.596							
	NT2RP400087	185.360					83.221	55.792	83. 494
								88.900	
	NT2RP400087							0.958	4. 164
	NT2RP400087							40.144	
	NT2RP400088								
55	NT2RP400089								
55	NT2RP400089	9 115.12	101.40	1110.21	11.44.70				

Table 114

			. 00 000 T	401 324 1	78.930	64.333	95, 484	61.641	85.667
	NT2RP4000902		188.808	401.324			0.372		0.771
	NT2RP4000906	0.305	1.603	0.000	0.000	0.170		0. 278	29, 737
5	NT2RP4000907	32.198	42.723	44. 472	21.830	25. 520	24. 599	19. 934	
•	NT2RP4000915	46.291	15.516	19. 755	9.014	17.749	25. 501	19.811	5. 257
	NT2RP4000916	16.757	34.708	57.738	23. 947	17.681	49.695	16. 463	25. 121
	NT2RP4000918	446.948	180. 459		104. 431	141.078	221.658	308.724	195. 950
	NT2RP4000925	33.696	20. 203	25. 426	9.727	8.694	5. 257	12. 183	6. 460
	NT2RP4000927	32.369	2. 391	13.088	3. 160	5.917	9. 429	12.242	9. 739
10	NT2RP4000928	132.499	77. 919	75.824	27.459	38.566	63.795	51.626	47.129
	NT2RP4000929	10.454	6. 358	16. 205	5.348	5.657	12.035	5. 522	3. 568
	NT2RP4000946	132.281	63.256	114. 387	25.969	53.023	57.751	42.531 0.000	26. 322
	NT2RP4000947	2. 292	0.165	0.000	0.681	0.000	0.000		0.000
	NT2RP4000949	61.713	79.888	67.197	17. 482	26.263	41.870	34.745 99.293	13. 260
	NT2RP4000955	138, 011	52.132	123. 547	28. 823	73.259	121.259	21.659	22.957
15	NT2RP4000959	41.008	45. 994	71.680	28. 437	32. 234	40.989		24. 213
.0	NT2RP4000962	18. 486	6.698	26.840	19.188	7.866	19.686	12.214	6.047
	NT2RP4000973	36.650	32. 445	36. 565	12.436	12.341	24.833	19. 337	14. 157
	NT2RP4000975	76.542	69. 291	152.889	24.672	28.007	28. 454	22.694	22. 187
	NT2RP4000979	34.880	19.409	37. 326	20.821	11.127	35. 561	8.305	14. 375
	NT2RP4000984	5. 549	5.330	0.000	9.035	5.964	4, 130	9. 900	5. 147
20	NT2RP4000986	67.644	33. 142	45. 802	10.889	17.544	33. 261	23.729	20.835
	NT2RP4000988	51.541	48.973	114.030	19.535	18.718	5.732	14. 224	16. 391 28. 297
	NT2RP4000989	59.625	24.400	48. 553	14.412	13.785	30.921	35.963 8.098	10. 761
	NT2RP4000990	18. 308	8.624	16.388	16.947	32.230	29. 187 47. 138	20, 380	42, 869
	NT2RP4000994	61.619	79. 591	73.376	19.693	19.056	48.697	18.081	61, 243
	NT2RP4000996	84. 850	105. 301	82.603	84. 356	51.465 34.967	41.069	18. 376	96, 597
25	NT2RP4000997	67.079	54.671	60. 172	11.766	11.811	15. 392	12.511	20. 370
_	NT2RP4001001	14.206	21. 359	18. 095 9. 361	5.116	9.588	16,002	13, 550	14. 012
	NT2RP4001004	33.229	16.130 32.280	76. 984	15.078	9. 382	26.487	11.510	24. 738
	NT2RP4001006	18.841	25.736	22. 167	10.117	15. 306	18, 272	18, 325	18, 908
	NT2RP4001009	66.828	26. 273	64. 129	11. 395	22.696	42,432	33.273	30. 440
	NT2RP4001010 NT2RP4001013	172.600	136.757	152.076	50, 579	71.395	91.790	74.989	69. 214
30	NT2RP4001029	51, 999	52, 569	51.080	19.391	11. 246	37.483	22.170	20, 460
	NT2RP4001029	50.398	28. 370	38. 461	20.941	14. 732	21.283	16.094	22, 458
	NT2RP4001041	63. 254	27. 315	44. 653	17,800	14.949	39, 536	29, 151	12.363
	NT2RP4001042	120.393	53. 507	99. 807	25.727	52.624	62.242	42.161	65, 349
	NT2RP4001046	84. 525	39.857	54.695	12.528	15.796	44,068	31.184	29. 152
	NT2RP4001050	23, 495	16.696	14, 229	3.130	7.595	15.142	37.084	15. 929
35	NT2RP4001051	55, 986	46.618	105. 231	34.838	19.098	22. 295	20.760	29. 183
	NT2RP4001057	106.673	52.182	65. 933	22.523	26. 382	66. 537	20.457	21. 945
	NT2RP4001063	170.235	69.039	102. 410	24.821	14.098	94.361	66.708	23.759
	NT2RP4001064	89. 983	57.290	64.770	15.070	12.139	42. 538	37.978	28. 126
	NT2RP4001067	32.210	18. 655	23. 175	7.147	6.320	18.181	17.994	8.877
	NT2RP4001078	70.346	22.808	30.478	9.119	13.915	11.118	32.316	11.554
40	NT2RP4001079	39.015	23. 923	38. 401	7.023	14.496	15.803	18.762	14. 515
	NT2RP4001080	14.552	29.116	54.653	6.580	5.732	7.627	7.008	8.413
	NT2RP4001086	62.838	43.770	64. 943	29, 980	22.792	56.125	30.073	45. 256
	NT2RP4001095	108, 108	110. 235	255. 542	37. 781	80.702	55.098	43.809	54. 938
	NT2RP4001098	70.282	49. 290	54. 985	17.657	20. 245	37.384	31.281	34. 153
	NT2RP4001100	197.231	163. 233	346, 289	64.078	75. 241	107.015	69.878	66.887
45	NT2RP4001105	230, 319	76.169	70. 257	26. 174	57. 028	86.626	87.810	59. 540 20. 912
	NT2RP4001110	57.855	44. 336	61. 199		18.898	18.716	33.736	
	NT2RP4001115	72.571	43.734	66.947	20. 426	27.358	20.977	23.470	23. 254 15. 958
	NT2RP4001117	53, 949	26. 454	27.949		12.786	27.164	36.880	32. 941
	NT2RP4001122	74.373	73.859	55. 273		24.494 17.045	39, 511 55, 106	52.069	29. 553
	NT2RP4001123	103.600	40. 395	69.670		48.688	55. 960	35.367	78.750
50	NT2RP4001126	70.020	118.846	92.913		4. 543	7. 932	6.088	3. 388
	NT2RP4001127	17.315	17. 921	20. 031			16.238	16.525	11.957
	NT2RP4001138	34. 858 89. 870	28. 363 104. 250	131.882			44,010	63.462	45. 180
	NT2RP4001143	10.496	8.968				2. 953	4. 275	13.549
	NT2RP4001148	121. 101	16. 961	36.641					17. 906
£	N12RP4001150	90. 570	29.463				28.002		17.678
55	MILKI 400 1130	1 30. 310	1 20. 400		1				

Table 115

		10 000 1	23.566	30. 231	13.969	15. 202	22.514	8. 474	15. 455
	NT2RP4001159	38.009		32.747	7. 435	8.821	8. 329	10.137	7.744
•	NT2RP4001162	26. 480	12.988			19.879	9. 183	5, 871	4.037
_	NT2RP4001170	22. 282	12.703	20.500	4.074			63.046	39. 356
5	NT2RP4001174	160.485	77.682	283.723	47.118	44.041	51.544		28.044
	NT2RP4001175	105, 636	84. 266	237.685	56. 987	37, 302	44.846	49.808	
	NT2RP4001176	316.295	539.044	440.109	306.340	44. 764	249. 181		321. 567
	NT2RP4001184	58. 252	23.348	36.224	15. 108	13. 298	29. 737	56. 984	15.700
	NT2RP4001198		120.100	81.937	37.566	13. 326	92. 551	80.670	61.997
	NT2RP4001199	22. 232	18.559	25.847	3.025	0.000	22.887	29. 205	23. 250
10	NT2RP4001205	167. 873	59.707	53. 222	31.978	27. 295	101. 042	75. 329	47. 196
		6.816	7.800	9, 463	4, 474	4.601	2.301	0.915	9. 232
	NT2RP4001207		9.826	9, 141	8, 107	1, 396	3.060	4. 469	2. 598
	NY2RP4001210	5. 482	21, 799	46.620	26.850	14, 691	14.012	16.268	14.828
	NT2RP4001213	18. 439		21.917	3, 759	2, 750	2.889	2. 203	1. 557
	NT2RP4001214	7.837	5.075		15. 168	7.172	11. 232	12.740	10. 296
15	NT2RP4001219	17. 372	12.922	29.465	22, 249	23. 349	41. 381	20.046	18. 506
	NT2RP4001228	60. 317	46.912	82.456		11.053	41.808	8, 307	26. 337
	HT2RP4001235	70.885	42.694	74.087	20.626		22.660	27.827	9, 288
	HT2RP4001256	53. 903	27. 494	40,975	9.302	9.044			16. 676
	NT2RP4001257	91.093	39. 253	66.828	12.871	33. 167	19.549	35.715	7. 635
	NT2RP4001260	30. 932	22. 193	31.916	6.755	16.733	19. 462	6.274	
	NT2RP4001261	203, 546	343.200	241.244	94. 907	116.433	194. 685	126.891	64.973
20	NT2RP4001274	29, 234	29.291	20.294	16.725	11.827	4.089	12.005	6.899
	NT2RP4001276	288. 394	86.186	155.256	76. 171	77. 526	99.724	126.975	37.044
	NT2RP4001283	602, 951	260, 199	332.966	68.876	287.262	624.729	534. 357	126.212
	NT2RP4001299	44.703	49. 576	35.736	19.564	12.675	15. 229	13, 741	18. 202
	NT2RP4001313	28.076	13.041	11.004	3, 551	7.304	11.207	9, 673	4.674
	NT2RP4001315	24.647	15. 443	17.362	12. 324	7.639	21.010	12. 223	11.809
25	NT2RP4001320	98. 164	61.534	65.437	15, 593	22.738	54.032	34. 155	23. 969
	NT2RP4001325	144.734	90, 080	132.401	61.000	64. 433	99.148	198.660	71. 382
	NT2RP4001336	33, 783	28.245	46.453	11,843	24.831	17.470	36.926	23.698
	NT2RP4001339	68. 525	15. 937	41.646	9.764	25.036	39.624	26. 253	9.570
	NT2RP4001343	161.856	91.193	100.371	27.738	38.512	92.415	57. 982	44.590
	MT2RP4001344	144, 107	58, 474	65.215	21. 137	22. 316	72. 157	71.543	28. 102
30	NT2RP4001345	50.445	32.733	43.703	11.121	15. 544	24.026	24. 553	13. 451
	NT2RP4001351	111.802	68, 455	97.136	54. 896	34. 425	45.604	34. 545	34. 491
	NT2RP4001353	19, 537	9.810	20.460	6.940	6.519	12. 325	7.907	7. 125
	HY2RP4001355	43.678	23. 203	33.304	7.482	15.675	24. 196	21.364	10. 692
	NT2RP4001367	14.283	17.653	14.776	4.211	8.006	2. 253	3.639	0.000
	NT2RP4001372	140, 185	27.600	56.900	12.537	24. 364	62.204	41.922	18. 450
35	NT2RP4001373	126.580	38. 189	93.856	23. 267	28. 220	77.754	42.832	38.641
	NT2RP4001375	62.861	32.389	48.017	13.250	23. 490	43.660	31.665	13. 296
	MT2RP4001379	77. 263	41, 191	123.636	24.440	18.057	56.629	33. 185	12.466
	NT2RP4001381	67.146	46.036	150.720	64.411	23.477	41.258	40. 245	17.295
	NT2RP4001386	47, 308	42. 524	147.963	19, 177	12, 559	15. 127	15.891	6.679
	NT2RP4001389	32, 461	38.092	48.803	17.637	14. 303	29.242	28. 109	24.013
40	NT2RP4001396	15.198	11.286	9.852	4.401	3. 270		5. 253	5. 075
	NT2RP4001407	13.731	19.546	21.832	9. 379	5. 846	11.131	8.899	4.678
	NT2RP4001409	26.965	45.073	26.488		6.075		11.306	7.105
	NT2RP4001410	111.952	58.388	89.502	31.596	42.948	111, 493	177.918	34.807
	NT2RP4001414	63, 484	72.860	54.366		26.471	40.346	21.075	42.279
	NT2RP4001424	18.505	15.050	18.180	8. 353	8.456		12. 261	8. 200
45	NT2RP4001433	28.627	47.828	111.176		3. 250		17. 950	7.176
43	NY2RP4001438	93, 429	51, 160	63.518	28. 266	34. 394		76. 382	59.077
	NT2RP4001442	45. 900	23. 169	80.514	5. 365	17.576			
	NT2RP4001447	20. 522	17.746						21.970
	NT2RP4001466	84.366	74.971		31. 341				
	NT2RP4001467	15. 268	25. 951	20.698	4.979				
50	NT2RP4001472	23, 447							
50	NT2RP4001474								
	NT2RP4001483								
	NT2RP4001488	27.970							
	MT2RP4001492								
	MT2RP4001498	25. 282	13.660	23.91					
	NT2RP4001502				8 60.785	58.64	7 81.80	46.693	100. 340
55									

Table 116

			-42	21 814	C 221 T	4 744	16 192 1	0 020 1	6.312
	NT2RP4001503	16.918	68.637	34. 943	6. 221	4.744	16. 123	9, 930	24.934
	NT2RP4001507	45. 444	50.856	165. 482	28.606	29. 404	30. 143	22.556	31.578
5	NT2RP4001510	32.998	28.050	63.008	35.045	3.511	13.039	13. 396	
-	NT2RP4001518	103.727	30. 191	54. 189	13.924	22.032	60.980	55. 131	21.835
	NT2RP4001520	99. 702	61.159	80.454	19.076	44. 823	57.892	65.886	85. 758 34. 713
	NT2RP4001523	74. 331	53.855	97.039	28.897	26. 233	31.769	22.342	
	NT2RP4001524	63.685	43.657	79.486	31.768	17.811	34. 268	61.096	32. 252
	NY2RP4001529	55.817	26.458	47. 156	18. 137	9.583	38.746	22.545	17.561
10	NT2RP4001531	76.425	49.034	79.547	19.985	15. 454	48. 895	27.165	35. 500
10	NT2RP4001546	475.672	254.067	158.609	114. 463	52.423	188. 321	90.884	193. 923
	NT2RP4001547	35. 657	46.341	75. 052	22. 751	21.180	18. 635	16.599	17. 284
	NT2RP4001551	15.709	5. 677	9. 034	3. 319	2.064	4. 065	8. 300	1.720
	NT2RP4001555	35. 187	13.947	15.040	6.049	8.613	14. 562	15.505	1.914
	NT2RP4001567	23.617	22.434	19. 944	10.030	13. 497	14. 121	17.021	12.931
15	NT2RP4001568	656.402	328.894	456.250	169.687	176. 926	432.308	269.108	137.575
10	NT2RP4001569	71.047	45.055	68. 921	13. 181	27. 919	55.014	36,067	22.875
	NT2RP4001571	31.048	30.838	25. 301	9.879	38. 867	28. 423	12.829	7.326
	NT2RP4001574	104.513	60. 846	51.480	12.719	37. 902	43. 358	52.975	26.473
	NT2RP4001575	99.868	54. 7921	66. 563	18.178	23.871	48.657	33,611	35.035
	NT2RP4001578	27.146	46. 286	41.253	12.060	16.868	28.516	38.747	21.566
20	NT2RP4001592	56.759	41.720	35.056	13.288	19.751	32.000	46.040	26.863
20	NT2RP4001593	34. 423	36. 251	40.059	19.801	27.006	22.857	28.378	30.708
	NT2RP4001605	35.830	55. 962	46.086	30.654	17. 304	12.782	25.954	20.171
	NT2RP4001606	36.059	22. 836	25.785	9.780	11.049	23. 731	22.906	11.246
	NT2RP4001607	12.252	38.564	26.768	11.976	11.793	10.856	12.358	17.689
	NT2RP4001610	41.606	26.761	24. 395	9. 284	13. 420	18.581	25. 355	17.897
05	NT2RP4001614	5. 120	1.451	3.713	3. 222	6.786	0.000	4. 236	3.006
25	NT2RP4001623	17.761	23.809	29. 296	18.722	11.464	7.465	7.749	11.940
	NT2RP4001626	39.777	77.553	31.850	125.728	14. 578	17. 234	15.665	43.780
	MT2RP4001634	42.268	33.465	29.710	15.079	5. 960	12. 998	22.448	22.801
	NT2RP4001638	28.002	28. 424	27.619	11.196	10.399	6.955	19. 293	11.952
	NT2RP4001644	13.937	31.012	33.018	11.442	10.696	15.844	17. 103	18.814
	NT2RP4001646	110.825	35.914	100.039	15.650	68.751	72.780	36.023	14.760
30	NT2RP4001656	113.964	57. 203	81.638	25. 444	41.071	67.708	57.712	34.629
	NT2RP4001666	75.518	31.622	54.757	17.666	17.943	29.002	29.742	13.617
	NT2RP4001670	143.248	64.754	95. 837	25. 903	20.467	58. 425	77.751	32.776
	NT2RP4001677	364.565	222.618	310.713	96. 394	105.468	224.860	256.793	96.732
	NT2RP4001679	225.706	136.839	407.981	82.012	82.799	62. 241	83.957	50.075
	NT2RP4001695	51.430	18. 839	33. 607	11.914	5. 205	20.014	20.606	3.263
35	NT2RP4001696	92.139	56.306	51.701	21.125	15. 829	67.642	34.335	27.080
	NT2RP4001699	20.126	24.412	12.024	6.153	9, 166	12.777	38.966	11.931
	NT2RP4001717	104. 794	22. 524	47. 196	16.831	10.332	44.003	26.697	10.303
	NT2RP4001719	4.115	3.996	6. 251	6.793	0.000	3.648	0.000	5.696
	NT2RP4001725	32.499	19.952	25. 192	14. 409	10. 172	27. 215	32.425	18. 951 19. 268
	NY2RP4001726	54. 527	36.453	64. 243	26. 169	28. 497	40.523	55. 394	4, 124
40	NT2RP4001730	12.704	4. 465	10.741	6.560	6. 940	4. 424	3.677 66.185	25.777
	NT2RP4001739	100.531	27.275	89. 269	26.597	21.415	57.785	33.008	41.898
	NT2RP4001741	110.382	99. 274	234. 294	44. 252	36.564	43.056 37.216	18.904	38.683
	NT2RP4001753	39.441	20. 491	71.424	37.461	1.805	5.657	5.650	2.825
	NT2RP4001760	14.764	11.531	4. 629		104, 482	137, 855	226.897	211.755
	NT2RP4001787	258. 392	145. 823	264. 342	128.018	19, 224	20. 959	21.785	26.319
45	NT2RP4001790	34. 934	24.033	47,502		29, 460	30. 950	41, 200	41.068
	NY2RP4001795	64. 250	59.518	90.887	55.846		11.542	8. 057	8.711
	NT2RP4001803	30. 124	17.002	33.008		5. 504	49.346	29.736	15.069
	NT2RP4001805	69.724	47.736	91.734		75. 066	114.744	91.752	13.588
	NT2RP4001809	249.052	50.599			13.677	59, 903	27. 216	21. 236
	MT2RP4001817	46. 954	36. 438				81.929		28. 953
50	NT2RP4001822	177.317	48. 258				10.465		6. 128
	NT2RP4001823	30. 502	15. 399				35, 276		20.301
	NY2RP4001827	65. 786	52. 243						63.747
	NY2RP4001828	265.068	110.898						18.265
	NT2RP4001836	136. 462	50. 159						
	NT2RP4001838								
55	NT2RP4001841	53. 995	81.543	68.608	23.556	51.873	33, 401	1 36.431	1 33.023

Table 117

	WYODANA A	147 467 1	17 445 [	38.764	4.795	19.911	39. 260	53. 938	8. 385
	NT2RP4001849	127.297	17.445		70.140				
	NT2RP4001861	247.889	113.986	152.565		77.706	119, 545	74. 993	93.651
5	NT2RP4001877	101.731	60.233	139, 463	63.735	37.564	43.408	50. 482	50, 974
	NT2RP4001879	52.547	46.318	81.300	25.097	20. 585	42, 533	33.249	30.904
	NT2RP4001889	70.569	45.627	140. 257	26.356	28. 442	18. 192	17.861	25. 113
	NT2RP4001893	25.380	22.592	43.017	18. 499	15. 138	9.424	8. 376	5.982
	NT2RP4001896	34, 081	20.051	44.749	10.547	15.271	19.037	14.839	13, 968
	NT2RP4001898	214, 122	125. 432	418.651	67.171	53.688	119,010	53.767	70.070
			53. 976	182. 276	39. 521	42.438	38.087	23.271	32, 169
10	NT2RP4001901	98.678				57.751	122.391	71.018	74. 327
	NT2RP4001910	37.857	50.894	99.896	25.518				19, 704
	NT2RP4001925	63.642	29.438	45.884	25.210	22. 129	45. 913	35. 236	
	MT2RP4001926	21,200	13.827	24. 573	7.083	11.581	7.544	10.754	9. 806
	NT2RP4001927	19, 268	17. 900	28. 017	9.329	12.222	11.234	14.514	7.786
	NT2RP4001931	97, 433	45.715	58. 255	21.472	23.167	20.695	41.852	23. 242
15	NT2RP4001933	94, 894	38. 536	49.118	18.868	30.123	17.978	20.041	20.718
,,	NT2RP4001938	286.138	121.070	279.936	37.391	35. 937	120.491	73.356	57.647
	NT2RP4001942	65.948	38. 369	38.848	27.689	31.221	62, 157	95,580	29. 143
	NT2RP4001945	41.368	18.714	27.898	8.014	14.644	17.772	15.860	11.677
	NT2RP4001946	26,736	25. 374,	44. 253	18.892	16. 137	18.739	15. 375	19. 575
				18. 880	3. 327	6.771	2.037	3.124	8. 202
	NT2RP4001947	3.902	6.862		20. 392	18.601	15. 837	9. 943	21. 246
20	NT2RP4001950	43.788	52.338	61.416					37. 538
	NT2RP4001953	74.594	54.521	201.576	35. 155	25. 200	19.900	24.690	
	NT2RP4001966	59.559	15.062	48.054	10.833	25.873	32.505	27.673	16. 883
	NT2RP4001970	250.998	97.493	91, 936	22.958	56. 420	113.696	71.723	47.051
	NT2RP4001975	66. 332	42.906	96.575	25. 793	35.690	65. 632	31.289	42.704
	NT2RP4001988	34.115	69.980	24. 419	10.144	8.048	24.865	25.619	34.649
25	NT2RP4001996	34.292	25. 552	27.656	12.286	17. 188	25.718	14.676	10. 275
25	NT2RP4002014	96.789	141.748	123.891	28.921	44. 195	55. 818	35.380	37.118
	NT2RP4002018	51, 913	24, 760	41, 235	10.014	19.687	23, 559	18.245	34.018
	NT2RP4002035	29.954	14. 435	25.087	12.863	31.601	28.211	23.642	22.189
	NT2RP4002043	22,692	22.569	32.501	19.499	11.381	22.102	20.695	23. 533
	NT2RP4002046	96.899	76. 132	55.715	18.254	26.488	53, 136	30.705	25.046
	NT2RP4002047	32.738	46.847	44. 327	32.723	15.068	26, 152	13.938	32, 144
30			18. 197	19. 425	11.638	8. 069	13. 935	10.066	12. 588
	NT2RP4002052	15.972				125. 176	113.563	77.593	83. 524
	NT2RP4002056	135.983	113. 302	169. 971	38.787			75.176	
	NT2RP4002057	84.885	34, 408	60.458	17.766	21.946	77.991		47. 433
	NT2RP4002058	23.685	18. 994	29.136	10.874	14.415	11.785	16.779	16.312
	NT2RP4002064	30.635	14.897	33.490	16.524	16. 922	12.258	15.014	25. 572
35	MT2RP4002071	44.464	31.989	67.896	26.934	14.700	35, 364	41.060	22.140
33	NT2RP4002075	12.341	23. 187	23.062	7.438	8. 387	13. 256	7.417	9. 609
	NT2RP4002078	29.846	42.027	82, 198	17.811	5. 991	41, 201	10.199	52. 455
	NT2RP4002081	188.987	84.568	105, 808	21. 123	35.926	97.846	71.564	35. 425
	NT2RP4002083	2.403	4, 985	0.000	0.314	0.000	0.000	1.168	0.000
	NT2RP4002099	78.239	28.086	39.672	11.893	30.439	25. 384	40.614	18. 182
	NT2RP4002106	58, 519	39.159	55.467	21. 121	12.917	56.398	104.992	30.602
40	NT2RP4002111	276.429	227.374	252, 398	129.656	67.040	206.459	208. 212	245. 585
	NT2RP4002112	24.864	26.459	24.698	12.961	9, 167	27.016	16.882	13.120
	NT2RP4002116	43.886	61.673	98, 270	42.933	38.005	36.286	25. 145	12.745
	NT2RP4002122	44,771	24, 552	24.373	26.789	12. 328	6.628	8.237	19.312
	NT2RP4002126	58. 138	23.058	51.469	13.176	13. 341	21.828	27.785	31.381
	NT2RP4002133	86. 426	80. 537	66.020	23.353	23.704	46.666	42.292	33, 380
45	NT2RP4002136	84.825	38.199	57.051	14.996	23. 918	31.464	46.186	26.328
45	NT2RP4002139				34.571	20.583	36.387	35, 415	37.823
		76.548	64.715	56.669	1 44 1 4 4	1 10 000	1	440	
	NT2RP4002174	100.223	26.805	135, 927	21.487	10.831	17.747	16.730	22.815
	NT2RP4002185	84.685	98.123	101.806	56.809	25.728	38. 576	50.054	42.202
	NT2RP4002186	76.426	104, 170	270.574	75.854	79.446	47.076	41.217	75.609
	NT2RP4002187	47.198	70.549	84, 418	12.734	27. 208	71.434	52.262	26.859
50	NT2RP4002188	35. 383	30. 278	67.328	48. 848	43,711	39. 200	18.696	45. 047
	NT2RP4002199	8.790	3.765	7.735	3.103	3.671	4, 856	6.602	4. 582
	NT2RP4002206	65.655	41.544	56.183	14. 975	16.172	23. 112	30.357	19.594
	NT2RP4002210	89, 632	39.449	49.442	26.733	14.817	29. 546	36.670	13.077
	NT2RP4002222	66.188	28.126	48.518	18. 433	9.476	18. 229	30.855	13.676
	NT2RP4002241	21.472	73.064	52.707	19.669	16.108	20.165	24.348	35.568
55	NT2RP4002248	89. 806	44.853	53.025	15. 207	28.490	47.016	40.320	26.933
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Table 118

	NT2RP4002250	9. 932	2.790	2.832	3. 032	2.884	3.939	3,541	2.731
	NT2RP4002259	98, 207	83.004	106.317	27.935	22.544	53.580	27.771	28. 361
	NY2RP4002268	76.648	38.869	74.529	23.758	30. 978	42.466	39, 855	19.465
5	NT2RP4002288	385.663	297.805	359.839	170.051	129. 643	303.550	199. 320	193.830
			20.072	47.837	6.799	18. 426	18. 201	11, 227	12.869
	NT2RP4002290	36.179	17. 225	18. 192	23, 131	9. 100	14. 492	16. 163	9.824
	NT2RP4002298	36.246			37.397	44.511	42. 955	43. 305	15. 782
	NT2RP4002308	106.632	73.744	244. 843					8. 392
	NT2RP4002308	32.611	5. 236	14. 575	3.239	0.000	18. 399	6. 762	
10	NT2RP4002336	58. 486	27.861	65.731	14.287	0.000	40.974	32.637	24. 822
•	NT2RP4002340	0.000	0,000	0.000	0.000	0.000	0.000	0.000	1.608
	NT 2RP4002361	58. 644	10. 427	47.735	8.516	9. 538	11.404	11. 251	4. 202
	NT2RP4002367	33.403	12.467	18.470	12.044	5.048	13.606	23.450	11. 551
	NT2RP4002368	30.961	37.918	39.910	11.210	13.572	15.090	26.947	17.073
	NT2RP4002377	54. 340	43.892	116.765	38.442	20.404	44.896	56.968	37.630
15	NT2RP4002408	13. 226	8.072	12. 192	7.437	5. 595	8.466	9. 233	1. 448
13	NT2RP4002425	9. 657	6. 220	8.381	3.685	1.438	1.029	3.019	3. 546
	MT2RP4002432	162.057	67.674	98.832	18. 405	27.254	50.612	54.723	48. 891
	NT2RP4002447	38. 164	33.834	62.023	30.863	14. 303	29. 507	8.767	8. 930
	NT2RP4002451	7.843	13.049,	15.746	6.677	3.617	8.815	1.747	9. 433
	NT2RP4002461	96.759	88, 219	116.998	47.479	57.340	32. 249	36.274	24.074
	NT2RP4002486	134.976	61.570	83.309	71. 309	46.898	61.095	41.576	21.740
20	NT2RP4002517	58.053	36, 106	59.653	12.934	25. 946	17.882	24. 902	21.801
	NT2RP4002556	43.020	59.649	60.047	47.543	16.113	30. 397	20.361	21. 390
	NT2RP4002569	55.960	24. 230	25.391	9.256	12.527	28.839	41.356	15. 537
	NT2RP4002587	66.993	24.539	29.137	12.319	10.000	27.896	26.210	13. 197
	NT2RP4002591	30. 924	17, 255	54.461	40.777	19.170	45.337	11,471	35.621
	NT2RP4002607	54.314	34. 936	46.019	25. 502	12.780	34. 916	29.754	19. 269
25	NT2RP4002627	77.997	65.880	94.854	27.581	43.756	52. 437	23.907	43.664
	NT2RP4002628	21.252	24. 628	31.576	38.351	13.833	13.934	20.421	19.758
	NT2RP4002630	70.308	48.663	165.068	28.270	26.685	23.920	21.463	25.050
	NT2RP4002639	34.573	25. 557	46.433	21.541	27.552	30.947	24, 555	20.118
	NT 2RP4002641	107.016	60.263	102.333	24.417	23.197	63. 371	31.978	22. 283
	NY2RP4002658	49.532	66.012	31, 405	43.805	11, 257	29. 226	40.300	36. 588
30	MT2RP4002669	139,676	37. 293	45, 595	14.602	38, 129	45.705	53.924	19. 231
	MT2RP4002677	20. 241	31.667	46.092	45.042	15, 952	20.098	16.586	53. 152
	NT2RP4002715	66.829	32.913	90.988	19.361	54, 330	49.099	48.752	33.038
	MT2RP4002750	74. 179	34. 932	56.851	17.150	20, 232	23.076	29.740	18. 218
	NT2RP4002784	67. 421	24.006	62.663	18.995	23.720	28. 427	58.514	16.510
	NT2RP4002791	28.944	34. 248	39.645	19.520	14, 437	25. 409	18.682	19.866
35	NT2RP4002811	191.101	48. 977	64.562	15, 450	18.301	79. 439	82.520	19.601
	NT2RP4002830	105.586	49, 177	76.222	25. 375	47.589	45. 374	21. 154	24.854
	NT2RP4002832	25.813	10,744	26.473	5. 157	5,007	10.239	3.522	4. 192
	NY2RP4002850	149.082	57.743	102.303	28. 532	37.913	75.770	47.566	29. 262
	NT2RP4002874	60.455	22.464	40.061	7.249	18.394	31.321	29.662	14.021
	NT2RP4002884	143.158	172,626	226.029	43.885	40.049	72.829	100, 195	80.578
40	NT2RP4002888	674.861	131,669	285. 125	53.073	130. 491	374.710	309.640	77.843
	MT2RP4002891	49. 251	19.998	83.408	45, 255	22.748	23.519	25. 198	32.282
	NT2RP4002894	52.025	17.730	44.439	15.465	30,670	53.933	19.786	17.490
	NT2RP4002896	62.611	29.872	36.349	8.160	14. 509	32.175	23.150	9.215
	NT2RP4002905	66. 278	20.133	27. 924	9.506	17. 387	37.876	9.098	11.326
	NT2RP4002907	133.109	146, 263	985. 435	80.359	7.640	119. 281	47.532	4.835
45	NT2RP5003459	104.697	52.694	23.001	28.403	58, 257	68.072	73.297	73.672
	NT2RP5003461	13, 597	25, 252	19,706	20.131	10, 924	4. 203	12.049	16.282
	NT2RP5003471	67.015	71.340	73.641	28.289	26,026	42.807	59.142	75.646
	NT2RP5003477	99.313	40.896	89.264		24, 470	47. 291	53.314	38.937
	NT2RP5003487	149, 480	394. 096	441.718		121.873	351.279	181, 435	545. 031
	NT2RP5003492	121.748	38. 219	55. 597	23.529	23. 174	61.042	47.790	32.333
£0	NT2RP5003500	28. 243	13.949			6.374	19, 974	10.924	7.373
50	NT2RP5003506	134. 622	138.997	142.784		50. 574	83. 904	56.185	74. 393
	NT2RP5003512	34.416	11. 927	16.738		7: 691	7.518	10.845	8.612
	MT2RP5003522	70.316	37.613			25. 328	22. 112	21.875	28. 029
	NT2RP5003524	37.812	24. 325			10.830	11.580	15, 958	15.290
	NT2RP5003527	548. 452	324. 151	547.100		240.483	435. 490	606, 993	396.820
	NT2RP5003531	218. 385	231.836	102.817		28. 603	161.069	24.666	55. 299
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Table 119

				Tabl	le 119				
	NT2RP5003534	52.710	32.028	38, 558	7.019	15.506	20. 156	15.510	8.979
					41.190	51.827	130, 501	91.960	29. 873
	NT2RP6000022	21.538	14. 233	20, 157	9, 477	8.940	8. 583	13.793	19. 548
5		71.839	29.419	34. 531	13.907	10.240	25. 335	35. 367	26.244
-	NT2RP6000050	64.066	28.604	49.917	15.400	35. 731	36. 275	41.783	27. 262
		158. 830	63.135	82.278	24.913	42.059	81, 152	81.658	52. 773
	NT2RP6000074	77.705	50.820	78.153	25.019	26.843	53.073	63.619	37.514
	NY28P6000100	50, 338	49.391	48.240	38.749	18.889	28. 023	21.991	17.677
	NT2RP6000123	93.881	40.481	91.240	14.231	12. 925	21. 554	11.762	21.172
10	NT2RP6000129	88. 985	47.556	65.182	18.250	6. 578	44. 353	35. 215	33. 928
	NT2RP6000147	32.349	57.944	378.808	14.768	31.975	24. 474	24.050	25. 120
	NT2RP6000163	25. 983	24, 930	19.397	3.670	5.619	4. 373	7. 249	6.228
	NT2RP6000181	156,005	46, 707	83.042	23.577	40.609	63. 752	82. 227	48. 686
	NT2RP6000182	88.398	83.770	188. 105	36.383	59.805	37. 752	28. 971	44. 514
	OVARC1000001	80.247	58.966	66.050	19.840	25.013	40. 518	55. 886	20. 561
15	OVARC1000003	20.948	26. 924	29, 257	6.511	9.613	12. 291	10.501	12. 527
	OVARC1000004	80.203	65.653	78.764	43.217	16.647	43, 045	49. 235	140. 643
	OVARC1000006	30.735	28.510	27.987	8, 280	13.556	26. 127	22. 928	6.489
	OVARC1000013	57.790	56.730	33.604	8. 558	12.487	25. 216	24. 982	13. 217 37. 444
	OVARC1000014	77.754	46. 427	51.294	14. 220	15. 288	23.140	45. 111	30. 588
	OVARC1000017	117.243	44, 469	63.710	20.838	20.603	56. 329	47. 142	92, 751
20	OVARC1000026	48.571	90.236	108.885	113. 202	47.802 28.515	62. 912 42. 393	36.018	23. 376
	DVARC1000035	49.364	29.380	53.296	24. 565	107.202	127. 890	99.068	110, 962
	OVARC1000037	217.386	150.962	123.831	103.776	55. 660	36. 598	18.334	26. 662
	OVARC1000058	126.770	102. 554	238.989 61.680	16.724	30.594	25.644	26. 946	50, 637
	OVARC1000060	69. 220	52. 141	21.327	11.889	11. 183	5. 229	8. 992	12.530
25	OVARC1000068	13. 131	13.623 75.808	53.487	27.968	38.426	35.714	36. 399	56. 941
25	DVARC1000069	101.314	24. 923	14, 847	12, 360	18, 401	4. 200	10. 123	10.463
	OVARC1000071	2485.301	555. 545	463.529	172.018	1135. 375	1656. 344	703.861	375.646
	OVARC1000073	28.000	26.575	47.619	26. 281	19.723	45. 186	32. 169	29. 140
	OVARC1000085	102.017	92.945	160,004	82.480	133.814	49. 366	47.840	68.991
	OVARC1000086	90. 269	76,669	42.857	41.659	25. 286	36.964	47.871	41. 838
30	OYARC1000087	19, 951	31.052	13.384	10.950	9.727	13.579	27.946	13. 255
	OVARC1000090	102.718	128. 317	77.866	86.960	52.554	50.597	59. 255	73.796
	OVARC1000091	20.738	22.588	16.835	15. 147	15.944	18.317	24.472	14. 038 30. 390
	OVARC1000092	45. 388	47.278	30.923	56. 969	21.795	27.471	41.724	47. 581
	OYARC1000105	56.618	51.625	28.040	39. 250	33.621	35. 440 45. 854	56.254	52. 554
	OVARC1000105	97.264	85.498	48. 102	31.853 28.577	38. 160	60.871	53.325	44. 146
35	OVARC1000109	114. 256	62.904	21.666	38. 682	21.936	24.745	30.026	25. 837
	OVARC1000113	34. 168	25.308 73.163	50.779	53. 005	19.962	35.869	27.590	39.625
	0YARC1000114	55. 942	9. 465	6, 445	3. 599	2.421	3.824	7.063	6.210
	OVARC1000133 OVARC1000137	41. 293	26.211	21. 220	13.873	15.408	25.975	29. 535	16.677
	DYARC1000139	84. 491	47.729	43. 252	31.553	15. 336	57.357	112.486	56.571
10	DYARC1000145	26. 915	13,800	13. 435	8. 493	4.736	16.675	21,905	9.604
40	DVARC1000148	95. 785	51.946	47.706	22.802	33.066	41.883	45. 597	39.685
	OVARC1000151	111.083	48.751	50.667	22. 177	24.840	81.184	53.839	31.939
	OVARC1000157	62.383	114.029	28.960		19.555	36.685	41.200	59.747
	OYARC1000162	5.118	14.000	6. 832		6. 337	4. 543	9.793	8. 590 49. 790
	OVARC1000168	81,607	75.614	57. 301		36.377	46.771	48.149	58. 589
45	OVARC1000169	78. 957	58.791	36.013		23.912	45. 597 64. 424		45.081
	OVARC1000178	106. 533	52.682	38. 529				9.722	7.079
	OVARC1000182	15.786	9.753					93.931	44. 381
	OVARC1000186	178.795							31,660
	OVARC1000188	55. 199							5. 421
	OVARC1000191	14.885							41.179
50	0VARC1000198								
	OVARC1000208					1			
	OVARC1000209 OVARC1000212								30. 436
	OVARC1000216						17.504		
	0VARC1000240								
	OVARC1000241					28.71	41.611	41.876	40, 107
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Table 120

	004001000040	61.518	33.009	33. 209	13.471	15.622	30.315	31.563	26. 366
	DVARC1000249		108. 103	77.039	42.563	81.235	81.095	78. 301	85. 308
	0VARC1000254	86.926			25.736	26. 168	36. 286	39.977	34, 354
5	0VARC1000255	60.970	39, 851	26.458				68, 179	55, 598
•	OVARC1000267	99. 396	106.106	72.814	56.946	76.696	67.094		
	DVARC1000275	1.361	3.837	0.000	0.676	1.682	4. 413	19.023	14.619
•	OVARC1000287	32.661	22.716	11.398	11.361	9.849	19.833	62.592	48. 262
	OVARC1000288	82.750	57. 876	28.088	21.493	23. 388	32. 508	34. 475	29.764
	OVARC1000298	23.487	30.867	16.778	9.152	10.710	22.218	12.148	15.140
	OVARC1000302	29.507	43.409	20.343	19.607	16.971	18. 175	10.089	15. 944
10	OVARC1000304	45. 645	44. 852	33.516	20.672	15.744	39. 549	33.592	42. 327
	DVARC1000307	24. 624	30.250	26.631	15, 444	18.919	21.450	27.043	23.654
	OVARC1000309	50.270	38.396	29. 381	16,928	23.152	40.904	33. 254	20. 287
	OVARC1000312	54. 891	39, 339	41.157	12.112	22.445	49.126	63.285	25.737
	OVARC1000313	62.108	49.417	23.463	16.503	20. 288	43.637	40.674	39.428
	OVARC1000321	38.317	91.534	39.988	16.691	58. 665	36.640	32.452	40, 394
15	OVARC1000326	58.790	34. 963	27. 371	25.834	24. 229	32.514	31.258	28.072
			45, 673	47, 401	27.601	25. 688	51.080	44. 339	24.826
	OVARC1000327	79.408			25.603	24. 396	42.607	52.669	29. 584
	OVARC1000331	67.541	33. 220	28. 427		12.062	15.090	16. 235	11. 984
	OVARC1000335	12.573	16.067	12. 457	10. 283		11.772	15. 807	14. 484
	OVARC1000347	10.404	19.839	9.744	14. 234	10.300		59. 346	36.658
20	OVARC1000348	104. 509	53. 231	29.087	28.611	27.286	49.055		
20	OVARC1800363	23. 207	29. 136	17. 234	17.138	22. 355	12.064	14. 282	17. 705
	OVARC1000377	24.447	20. 967	8.919	11.225	9.000	9. 306	12.677	10. 839
	OVARC1000382	43. 425	38. 484	25. 520	12.983	19. 971	27.581	24.011	20.004
	OVARC1000384	39.526	33.430	34.510	29.733	34. 546	26. 194	23.240	27.218
	OVARC1000401	19.377	21.365	10.833	14.856	8. 159	13.368	15.387	15. 593
	OVARC1000406	246.308	104.316	212.801	47.902	275. 450	229. 284	231.727	63.004
25	OVARC1000407	37.707	28.148	15. 167	29.769	18. 198	20. 301	24.339	24. 226
	OYARC1000408	176.546	182.488	168.003	92. 253	152.822	131.022	104.696	123.181
	OVARC1000410	132.351	71.592	33.987	19.006	47. 593	63. 597	105.036	45.064
	OVARC1000411	24.928	45.964	21.466	16.795	18.354	16.759	17.621	24. 921
	OVARC1000414	53.052	80. 288	77.929	45.828	64. 588	36.694	43.527	34.813
	OVARC1000420	210.281	97.795	116.314	65.770	48. 502	138, 372	122.961	79.364
30	OVARC1000421	126.414	65. 308	43.609	41.965	30.984	66.717	77.617	43.013
	OVARC1000427	85. 522	76.052	61.132	54.694	43. 202	55. 414	85.904	58. 425
	0VARC1000431	29.754	43. 257	31.464	59.910	40. 269	33, 174	24.118	40.748
	OVARC1000437	101.746	108.759	36.433	33.368	38,706	67.360	68.627	59.999
	OVARC1000439	55. 100	39. 820	23.665	17.682	26.837	27.173	37.589	27.642
	0VARC1000440	9. 304	16.390	4.607	5.910	7.569	12.799	5.759	8.778
35	0VARC1000442	71.954	97. 290	60.169	48.043	50. 925	49.780	49. 452	49.132
00	DYARC1000443	23.336	24.854	21.466	6.313	14. 231	15.300	17.929	17.277
	OVARC1000461	38.961	27. 338	30.933	18.801	25. 228	29.577	31.675	33.815
	OVARC1000465	24. 244	26. 635	23. 588	15.988	16. 431	17. 245	18.033	20.237
			45. 309	35. 183	22.710	29. 028	42.270	78. 325	34.551
	0VARC1000466	78.845	41.545	26.636	17.995	24. 535	32.636	50. 520	33. 453
	OVARC1000467	68. 457		34. 473	51.974	38.874	30. 248	35. 482	44.070
40	OVARC1000470	79.505	56.390				53.144	60.639	36.861
	OVARC1000473	104. 826	46.950	38.060	19.545	49.878	11, 592	14. 222	14.645
	OVARC1000479	13.043	22.838	18. 446		14. 611		32.384	37.475
	0VARC1000484	81.135	119,477	61.550	71.199	61.618	42. 186	12.533	17. 483
	OVARC1000486	43.060	37.552	15. 873	26.931	21.970	20.014		
	OVARC1000496_	6.894	5. 795	2.024	9.550	5. 845	4. 482	5. 597	6.952
45	OVARC1000520	10. 944	13. 261	5. 969	10.975	8.640	4. 581	5.177	7.377
	OVARC1000522	57.377	36, 524	49. 921	34. 183	62.162	27.574	36.847	42.071
	OVARC1000526	89.641	108.239	58.125	65.691	63.235	45. 200	63.148	66.145
	OVARC1000529	57.424	54.050	21.682	25.091	30.072	29. 592	53.851	44.743
	OVARC1000533	259.058	92. 210	92. 325	54.816	108, 661	158. 123	180.752	58.313
	OVARC1000543	9.147	20.003	8. 468	14.598	10.808	7.160	7.656	9.778
50	OVARC1000550	51.120	35.681	19.454	19.769	24. 341	23.780	29.758	19.911
			109.455	53.476	65.549	62.372	43.061	54.040	60.551
50	OVARC1000553	106.477	103.433						1 AA 417
30	OVARC1000553 OVARC1000556	84.636	47. 645	29. 302	13.010	36, 300	34.811	56.871	26.716
50	OVARC1000556	84.636		29. 302 15. 138	13.010 26.106	36.300 16.920	12. 137	18.572	18.092
30	0VARC1000556 0VARC1000557	84. 636 30. 381	47.645						
30	OVARC1000556 OVARC1000557 OVARC1000561	84.636 30.381 130.212	47.645 33.997 131.086	15.138 62.529	26, 106	16.920	12. 137	18.572	18.092
55	0VARC1000556 0VARC1000557	84. 636 30. 381	47.645 33.997	15, 138	26.106 70.306	16. 920 56. 212	12. 137 50. 615	18.572 65.315	18.092 60.071

Table 121

					FOS T	es eex	100 010 1	101 660	126.973
	OVARC1000576	322.369	178.635	134.897	63.583	51.558			
	OVARC1000578	41. 245	47.399	27.512	62. 221	23.000	19. 402	20.991	28. 222
	OVARC1000581	19.381	18.054	16. 597	12.946	10.926	16.921	23. 587	17.958
5	OVARC1000586	58,760	84.513	39.858	53. 327	17.530	41.985	63.279	95.673
	OVARC1000588	52.736	46.547	28.747	35, 144	19.236	20. 189	27. 881	28. 239
	DVARC1000505		21.584	16.038	16,026	12,949	28.632	20. 949	12.415
		25.011			146.619	111.039	103, 900	84. 581	117.758
	DVARC1000622	236.401	229.625			23.116	39, 517	52.845	27.699
	OVARC1000636	62.041	58.870	30.872	26.680				24.570
	OVARC1000540	37.774	40.454	27.435	25. 421	14, 327	20.971	27. 326	
10	OVARC1000549	119.925	80.531	59. 932	34. 951	42.653	66.545	126. 333	64.422
	OVARC1000661	91.942	47,731	46.674	29.765	29.826	53. 562	68.611	41.478
	OVARC1000677	47.303	42.727	39.478	18.654	17, 990	29.788	33. 925	31, 139
				32.060	37.092	23. 552	26. 846	42.330	32.378
	OVARC1000678	53.878	40.134		13.826	12.729	13. 248	18. 589	22. 125
	OVARC1000679	25. 552	33.892	27.236		25, 157	35.864	32.183	28. 963
	OYARC1000681	64.996	39, 676	33.010	23.036				
15	OVARC1000682	89. 453	46.031	48.073	26. 181	22.664	56.539	67.656	36. 205
	OVARC1000689	40.786	43, 141	31.489	15.450	18.494	36.522	52.050	30. 362
	OVARC1000700	55. 561	65.260	46.443	51.382	36,724	40.865	31.889	43.299
	DVARC1000703	68. 421	67.574	44.166	43. 328	32.848	43.707	34.063	33.710
	OVARC1000722	90.588	55. 674	40.426	28.083	33.617	39.059	84.669	53. 295
				64. 375	36.671	45.678	62.745	120, 014	59.080
00	OVARC1000726	223, 039	61.254			24.747	45, 216	39. 732	28. 241
20	DVARC1000727	101.498	52.857	32.778	21.030			24. 052	21. 653
	OVARC1000730	32.092	36. 451	14. 144	26.825	11.752	14.326		
	OVARC1000741	93.409	\$2.169	37.001	21.498	22.633	47.358	43.609	24. 156
	OVARC1000748	18.880	20.011	11.250	10.152	10.039	12.336	11.833	14. 185
	OVARC1000764	94.412	66.494	49. 103	37. 950	38. 405	\$7.102	51.799	45.024
	OVARC1000769	61.249	87. 994	63.412	48.573	49.372	45.621	37.596	54.747
25	DVARC1000771	17.704	22.392	12.731	11.680	15.094	14, 537	11.734	13. 386
23	OYARC1000773	309, 712	63.691	128.640	93. 505	135.643	247.891	47.762	56. 423
		39. 822	40.473	19.087	17, 945	17.047	20. 931	22.217	24. 299
	OVARC1000775		40. 229	23.354	27.887	19,703	25, 351	15. 434	16.858
	OVARC1000778	57.819			4.604	3.376	6. 590	5, 881	4.469
	OVARC1000779	13.359	9.700	3.596			18. 289	13.313	9. 987
	OVARC1000781	28. 426	18. 324	19.364	8.066	2.533			
30	OVARC1000787	57.756	46.552	31.435	36. 327	24.660	31.315	26. 423	26.916
	OVARC1000789	\$6.045	42.830	21.343	32. 278	26. 180	29.815	35. 483	31.401
	OVARC1000800	152.906	115.192	91.456	100.625	80.665	74.709	72.586	83. 426
	OVARC1000802	56.307	41.592	29.261	21.865	29.614	38.004	29. 144	37.338
	07ARC1000810	117, 305	73.073	45. 217	47.024	30.840	54, 331	22.585	30.212
	OYARC1000811	24. 376	21.125	12. 822	10,066	8.476	14.818	12. 129	13.407
		109.717	173.696	116.374	110, 400	99.820	81.598	52.542	70.043
35	OVARC1000814			29, 109	10.508	18.910	26.961	43. 388	30. 931
	OVARC1000816	38.942	32.627		4. 922	1. 435	3.770	5. 107	7.052
	OVARC1000817	7. 152	7.754	5. 073		27.698	43.601	43. 333	33.619
	OVARC1000834	52.593	59.148	30.623	25.871			47. 279	73. 330
	OVARC1000846	128.045	121.550	80. 555	82.014	63.814	79.270		
	OVARC1000850	63.194	47.834	24. 998	22. 731	23.832	31.759	43.348	29.789
40	OVARC1000853	47.482	127.726	57, 523	25. 169	55.048	41.556	32.136	37.576
40	OVARC1000862	31, 255	26.218	21.640	13.240	25.873	16.507	16. 932	8.079
	OVARC1000873	59.654	49.105	31.649	32. 533	37.513	39.866	44. 461	30. 226
	OVARC1000875	178.627	94. 134	92.359	64.818	79.244	116.581	163. 150	75.514
	OVARC1000876	8.798	15.017	5. 566	12.799	6.112	8, 158	8. 444	16.825
	OVARC1000883	44. 435	33.208	17, 857	13, 562	21.585	25. 327	28.768	27.716
			16. 263	7. 277		8.434	58.765	18, 303	13.712
45	OVARC1000885	11.029					10.692	35. 601	21.522
-	OVARC1000886	41.813	40.085	18.851			1	128.741	96. 438
	DVARC1000890	216.895	167.860	92. 458	00.405				
	OVARC1000891	20.905	24.028						12.875
	OVARC1000897	9.048	31.172	6. 976			7. 384		9. 271
	OVARC1000912	15.809	11. 325		14, 551				
	OVARC1000914		35. 138				18, 401	14, 325	19. 336
50	OVARC1000915						40.708	37.700	37.607
	OVARC1000916								29. 531
	OVARC1000924								
	OVARC1000928								
	OVARC1000936								
	OVARC1000937	50.958	48. 239	37. 55	26.648	23.630	35.710	( 31.349	1 33.003
55									

Table 122

•			1 au	16 122				
OVARC1000945	72.670	66.756	35.734	31.061	28. 439	44. 288	57. 299	34.609
	13, 138	9.821	6.873	5.701	6. 145	7. 947	8. 485	6.560
	53. 521	35.128	27.412	25.007	31.512	30.356	47.794	38.003
	73.657	56.906	34. 594	53.936	29.777	37.237	43.699	40.734
		304.478	264.514	101.674	301.925	170. 334		211. 921
	09. 457	89.334	92.736	42.962	107. 425	66.304		104. 440
OVARC1000971	23.347	22.555	11.767	9.454	10. 751	11.968	14. 346	9. 949
DVARC1000975	38.653	41.668	22. 926	16.702	21.947	23,016	30. 329	22. 999
OVARC1000976	5. 549	11.344	5, 097	7.562	5. 992	4, 915	5.760	8. 357
OVARC1000981	38.051	38.818	23.473	34. 246	24. 179	25. 155	27.878	49. 594
OVARC1000982	18. 237	20.180	8.868	15. 397	12.870	12.622	17.681 40.246	16.489 32.960
DYARC1000984	64, 280	32.461	21.258	21.860	23. 534	26.715	46.967	67. 398
DVARC1000995	98.670	98.801	50. 363	66. 552	60.125	43.967	20. 244	19, 657
OVARC1000996	23. 461	22.409	9. 648	11.387	13.424	14. 277 72. 322	63.450	71.881
OYARC1000999	142.766	147.956	91. 391	112.389	86. 587	96.923	91, 581	110. 353
	196.742	223.698		137.198	124.411	5, 392	7,712	7.076
OVARC1001004	15.837	24.777	8. 416	6.761	11.301	9, 756	20,656	11. 492
DVARC1001010	20.746	21.844	10. 176	13.214	12. 974 29. 627	26.821	32.007	29.717
OVARC1001011	56. 262	49.134	31.219	40. 269	481. 589	213. 259	236.252	156.604
***************************************	267.698	257.417	369.890	123.083	17, 403	12.728	21.746	22.424
OVARC1001032	25.684	32.175	13.978 18.682	17.255	24. 225	14. 437	22.093	19.148
OVARC1001034	26.408	30, 129	24, 957	24.612	28.412	29.918	36, 871	30.300
OVARC1001038	38.346	41.992	57.680	96.342	37.120	36.870	51,690	65.780
0VARC1001040	98.109	176.563	45, 646	73.484	59. 177	42, 401	48. 353	76.436
DVARC1001041	93. 629 29. 011	33.627	14. 802	21.262	17.318	16.763	22.227	22.829
OVARC1001044 OVARC1001049	156.011	131,461	99,014	60.845	95. 518	95. 243	124. 468	83.710
OVARC1001049	180.769	195.784	75.946	127.551	72.219	104.988	166.021	161.466
OVARC1001054	44.196	25.475	14, 270	15.193	14.800	17. 493	25.623	19.511
OVARC1001055	49.946	52.425	26.074	16.256	16.038	22.736	26. 492	26.988
OVARC1001062	9.764	52,550	13.991	22.860	14. 380	12.344	7.304	17. 143
OVARC1001065	20.300	19.807	20. 195	9,804	10.947	15. 910	27.631	19. 975
0YARC1001068	56.993	44.653	31.867	20.677	17. 254	28.843	44. 829	31.704
OVARC1001072	156.343	67.114	52.898	30.164	30.884	59.064	66.747	40. 238
DVARC1001073	34.815	40.406	29.440	33, 203	20.617	29.525	38, 538 22, 029	15.581
OVARC1001074	18.735	18.807	6. 927	9.591	8. 229	12.569	97. 102	51.664
OVARC1001078	170.789	81.144	63. 392	42.879	16.558	51.666	25. 272	24.844
OVARC1001085	48. 583	37.562	22.446	18.020	19.838	42.555	39.613	26.858
OVARC1001086	94, 509	38, 291	23.565 39.117	31.558	15.085	41.655	65, 548	38.043
OVARC1001091	59.024	54. 767 48. 386	35. 270		27.135	48.099	68. 542	28. 251
DVARC1001092	78.369 9.822	12.079	8. 053			6.895	13.769	8.849
OVARC1001104 OVARC1001107	132.584	59.642	57.112		46.497	103.685	120.752	61.479
OVARC1001113	35, 730	35.073	29.872			24.132	39. 291	35.356
OVARC1001117	91.761	65.878			23.367	45.042	42. 492	38. 455
OVARC1001118	78.150	72.874	_			49.123	35. 261	47.146
OVARC1001125	19.282	29. 524					19.586	21.569
OVARC1001129	26.932	18.396					19.030	7.081
	7.132	10.388	7.883					8. 385 99. 862
OVARC1001132				77.068	99.486	165, 174		
OVARC1001132	308.799	242.318					20 000	24 41
OVARC1001138 OVARC1001141	308.799 48.972	28. 503	23. 912	13.741	19. 193	23.582		
OVARC1001138 OVARC1001141 OVARC1001154	308.799 48.972 66.885	28.503 91.460	23.912 43.947	13.741 37.042	19. 193 36. 702	23. 582 48. 431	80.339	79.15
OVARC1001138 OVARC1001141 OVARC1001154 OVARC1001161	308.799 48.972 65.885 71.634	28, 503 91, 460 56, 342	23. 912 43. 947 31. 340	13.741 37.042 42.482	19. 193 36. 702 14. 597	23. 582 48. 431 25. 244	80.339 28.686	79.15 26.64
OVARC1001138 OVARC1001141 OVARC1001154 OVARC1001161 OVARC1001162	308. 799 48. 972 66. 885 71. 634 80. 697	28.503 91.460 56.342 81.514	23. 912 43. 947 31. 340 58. 69	13.741 37.042 42.482 43.494	19. 193 2 36. 702 2 14. 597 1 34. 028	23. 582 48. 431 25. 244 46. 796	80, 339 28, 686 40, 262	79.16 26.64 50.82
OVARCIOO1138 OVARCIOO1141 OVARCIOO1154 OVARCIOO1161 OVARCIOO1162 OVARCIOO1163	308. 799 48. 972 66. 885 71. 634 80. 697 170. 857	28. 503 91. 460 56. 342 81. 514 43. 068	23. 912 43. 947 31. 340 58. 69 59. 42	13.741 37.042 42.482 43.494 4 17.764	19. 193 2 36. 702 2 14. 597 4 34. 028 4 29. 285	23.582 48.431 25.244 46.796	80, 339 28, 686 40, 262 90, 481	79. 16. 26. 64 50. 82 55. 48
OVARC1001138 OVARC1001141 OVARC1001154 OVARC1001161 OVARC1001162 OVARC1001163 OVARC1001167	308.799 48.972 66.885 71.634 80.697 170.857 77.273	28. 503 91. 460 56. 342 81. 514 43. 068 85. 145	23. 912 43. 947 31. 340 58. 69 59. 424 46. 744	13.741 7 37.042 0 42.482 7 43.494 4 17.754 5 44.761	19. 193 2 36. 702 2 14. 597 4 34. 028 4 29. 285 8 32. 264	23. 582 48. 431 25. 244 46. 796 91. 606 35. 631	80, 339 28, 686 40, 262 90, 481 32, 410	79, 16 26, 64 50, 82 55, 48 29, 95
OVARC1001138 OVARC1001141 OVARC1001154 OVARC1001161 OVARC1001162 OVARC1001163 OVARC1001167 OVARC1001167	308.799 48.972 66.885 71.634 80.697 170.857 77.273	28. 503 91. 460 56. 342 81. 514 43. 068 85. 145 )5. 674	23. 912 43. 947 31. 346 58. 69 59. 424 46. 741 9. 30	13.74) 7 37.042 9 42.482 7 43.494 17.764 6 44.760 2 5.67	19. 193 2 36. 702 2 14. 597 4 34. 028 4 29. 283 8 32. 264 4 5. 124	23. 582 48. 431 25. 244 46. 796 91. 606 35. 631 9. 510	80. 339 28. 686 40. 262 90. 481 32. 410	79. 16 26. 64 50. 82 55. 48 29. 95 9. 74
OVARC1001138 OVARC1001141 OVARC1001154 OVARC1001161 OVARC1001162 OVARC1001163 OVARC1001167 OVARC1001169 OVARC1001170	308.799 48.972 65.885 71.634 80.697 170.857 77.273 10.634 48.257	28. 503 91. 460 56. 342 81. 514 43. 068 85. 145 75. 674 49. 203	23. 912 43. 947 31. 346 58. 69 59. 42 46. 74 9. 30 32. 87	13.741 7 37.042 9 42.482 7 43.494 4 17.754 5 44.761 2 5.674 9 28.36	19. 193 2 36. 702 2 14. 597 4 34. 028 4 29. 283 3 32. 264 4 5. 124 6 23. 146	23. 582 48. 431 25. 244 46. 796 91. 606 35. 631 9. 510 21. 439	80, 339 28, 686 40, 262 90, 481 32, 410 12, 220 43, 645	79.16 26.64 50.82 55.48 29.95 9.74 39.07
OVARC1001138  OVARC1001141  OVARC1001154  OVARC1001162  OVARC1001163  OVARC1001167  DVARC1001169  OVARC1001171	308. 799 48. 972 66. 885 71. 634 80. 697 170. 857 77. 273 10. 634 48. 257 71. 425	28. 503 91. 460 56. 342 81. 514 43. 068 85. 145 75. 674 49. 203 65. 034	23.912 43.947 31.346 58.69 59.42 46.740 1 9.30 32.87 5 38.59	13.741 7 37.042 9 42.482 7 43.494 4 17.754 5 44.761 2 5.67 9 28.36 5 39.74	19. 193 2 36. 702 2 14. 597 4 34. 028 4 29. 285 8 32. 264 4 5. 124 6 23. 146 6 29. 129	23. 582 48. 431 25. 244 46. 796 91. 606 35. 631 9. 510 21. 439 40. 964	80, 339 28, 686 40, 262 90, 481 32, 410 12, 220 43, 645 39, 089	79. 15. 26. 64. 50. 82. 55. 48. 29. 95. 9. 74. 39. 07.
OVARC1001138  OVARC1001141  OVARC1001154  OVARC1001162  OVARC1001163  OVARC1001167  DVARC1001169  OVARC1001171  OVARC1001171	308. 799 48. 972 65. 885 71. 634 80. 697 170. 857 170. 857 77. 273 10. 634 48. 257 71. 425	28. 503 91. 460 56. 342 81. 514 43. 068 85. 145 75. 674 49. 203 65. 034	23. 912 43. 947 31. 340 58. 69 59. 42 46. 740 9. 30 32. 87 5 38. 59 2 67. 40	13.741 7 37.042 9 42.482 7 43.494 4 17.75 5 44.761 2 5.67 9 28.361 5 39.74 6 103.30	19.193 2.36.702 2.14.597 4.34.028 4.29.285 8.32.264 4.5.124 6.23.146 6.29.125 7.65.935	23. 582 48. 431 25. 244 46. 796 91. 606 35. 631 9. 510 21. 433 340. 964 360. 125	80, 339 28, 686 40, 262 90, 481 32, 410 12, 220 43, 645 39, 089 54, 280	79. 16/ 26. 64/ 50. 82/ 55. 48/ 29. 95/ 9. 74/ 39. 07/ 54. 22/ 60. 38/
OVARCIO01138 OVARCIO01141 OVARCIO01154 OVARCIO01163 OVARCIO01163 OVARCIO01167 DVARCIO01169 OVARCIO01170 OVARCIO01171 OVARCIO01173 OVARCIO01173	308. 799 48. 972 65. 885 71. 634 80. 697 170. 857 170. 857 77. 273 10. 634 48. 257 71. 425 116. 007 245. 124	28. 503 91. 460 56. 342 81. 514 43. 088 85. 145 75. 67 49. 203 65. 034 101. 332 107. 904	23. 912 43. 947 31. 340 58. 69 59. 42 46. 74 9. 30 32. 87 5 38. 59 2 67. 40 8 82. 42	13.741 7 37.042 9 42.482 7 43.494 4 17.75 5 44.76 2 5.67 9 28.36 5 39.74 6 103.30 1 85.01	19.193 2 36.702 2 14.597 4 34.028 4 29.283 5 32.264 5 124 6 29.125 7 65.933 4 77.97	23.582 48.431 25.244 46.796 91.606 135.631 9.510 140.964 140.964 145.455	80, 339 28, 686 40, 262 90, 481 32, 410 12, 220 43, 645 139, 089 54, 280 105, 359	79. 16/ 26. 64/ 50. 82/ 55. 48/ 29. 95/ 9. 74/ 39. 07/ 54, 22/ 60. 38/ 82. 55/
OVARC1001138 OVARC1001141 OVARC1001154 OVARC1001162 OVARC1001163 OVARC1001167 OVARC1001170 OVARC1001171 OVARC1001173 OVARC1001173 OVARC1001176 OVARC1001176	308.799 48.972 66.885 71.634 80.697 170.857 77.273 10.634 48.257 71.425 116.007 245.124	28. 503 91. 480 56. 342 81. 514 43. 068 85. 145 75. 674 49. 203 65. 039 101. 332 107. 900 157. 050	23.912 43.941 31.346 58.693 59.422 46.746 3 9.30 32.87 5 38.59 2 67.40 3 82.42 5 72.13	13.741 7 37.042 9 42.482 7 43.494 4 17.76- 6 44.765 2 5.67- 9 28.36 5 39.74- 6 103.30 1 85.01 6 68.29	19.193 2.36.702 2.14.597 4.34.028 4.29.285 4.29.285 6.23.144 6.29.125 7.65.933 4.77.970 0.69.36	23.582 48.431 25.244 46.796 91.606 21.435 35.631 9.510 521.435 340.964 360.125 5145.455 772.29	80, 339 28, 686 40, 262 90, 481 32, 410 0 12, 220 0 43, 645 4 39, 089 0 54, 280 1 105, 359 0 67, 658	79. 152 26. 644 50. 825 55. 486 29. 956 9. 74 39. 07 54, 22 60. 38 82. 55
OVARCIO01138 OVARCIO01141 OVARCIO01154 OVARCIO01163 OVARCIO01163 OVARCIO01167 DVARCIO01169 OVARCIO01170 OVARCIO01171 OVARCIO01173 OVARCIO01173	308. 799 48. 972 65. 885 71. 634 80. 697 170. 857 170. 857 77. 273 10. 634 48. 257 71. 425 116. 007 245. 124	28. 503 91. 460 56. 342 81. 514 43. 668 85. 145 15. 67 49. 203 65. 033 101. 337 107. 900 157. 056 349. 53	23.912 43.941 31.346 58.69 59.42 46.741 9.30 32.87 57.40 38.59 67.40 382.42 572.13 8 32.80	13.741 7 37.042 9 42.482 7 43.494 4 17.765 5 44.761 2 5.67. 9 28.36 5 39.74 6 103.30 1 85.01 6 58.29 4 26.68	19.193 2 36.702 2 14.597 4 34.028 4 29.283 8 32.264 4 5.122 6 23.144 6 29.123 6 6 65.933 4 77.970 0 69.363 3 20.34	23.582 48.431 25.244 46.796 91.606 35.631 4 9.510 5 21.433 6 40.964 6 60.123 7 72.297 8 25.531	80, 339 28, 686 40, 262 90, 481 32, 410 112, 220 43, 645 43, 645 44, 645 45, 645 46, 645 47, 645 48, 6	79. 161 26. 641 50. 82: 55. 48: 29. 95: 9. 74 39. 07 54. 22 60. 38 82. 55 67. 80

Table 123

									65 A16
{	OVARC1001202	122.810	79.160		45. 179	34, 305	57.752	54. 521	52.075
	QVARC1001206	42.615	25, 397	25. 932	13. 326	28, 104	29.089	32.918	22.690
	OYARC1001209	72.876	58.366	36.700	24. 151	40.859	38, 440	59, 191	47, 601
			13, 311	13.625	12.687	15, 459	16, 636	29.651	23, 801
	OVARC1001219	33.632				10. 225	25. 581	20.058	17.564
	OVARC1001222	32.786	21.648	10.686	9.886				
	OVARC1001232	117.540	87.613	50.146	34.554	30. 246	57.933	49. 208	37.950
	OVARC1001240	75, 374	60,625	38.831	32.204	26. 238	32.631	20. 938	29. 225
	OVARC1001243	9, 543	16.485	6. 223	5.619	1.978	7.592	11.212	9. 204
			111.321	69.720	46. 121	39. 223	93, 281	105. 487	89.348
	OVARC1001244	169.003			59. 295	307.379	168. 939	66.384	180.606
	OVARC1001246	102.652	232.219		17.972	24. 516	29, 579	38, 406	32.633
	OVARC1001247	51, 814	49. 398	25. 400					
	OVARC1001260	53. 551	100.419	29. 364	25.020	34.864	30. 489	28. 556	34. 131
	OVARC1001261	48. 536	42.267	28. 153	13.070	26, 118	36. 541	37.660	22.612
	OVARC1001268	51,904	118.717	47. 463	24.361	63.661	38. 492	51, 108	43. 123
	OVARC1001270	20.955	18.655	11.209	10.629	7, 297	10.404	10,615	9, 730
				59. 789	67.369	40, 952	49.040	49. 902	56.550
	OVARC1001271	82.087	105.253				4.070	1.658	4, 939
	OVARC1001282	2. 151	7.862	2.074	5, 144	2.146			
	OVARC1001296	11.865	15. 267	7.897	10.844	6. 153	11.518	15, 515	10. 296
	OVARC1001306	25. 532	50.725	28.628	24.049	17.847	22.716	24. 404	32. 492
	OVARC1001314	12.995	19.789	11, 346	14. 481	11.454	16.041	17.642	15. 122
	OVARCIDO1316	14. 093	43.453	9. 049	9. 287	10.402	12.676	9. 571	8. 634
			224. 291		140. 553	147. 173	134.506	88. 940	124. 623
	OYARC1001329	236.298					18.781	21.774	14. 306
	OVARC1001330	34.063	30.737	21. 299	12.416	9, 409			
	OVARC1001336	64. 433	86.449	37, 979	30.312	22. 554	34. 649	46, 151	35. 127
	OVARC1001338	29.434	27.732	16. 123	16.132	16. 945	20. 146	25. 217	26, 946
	OVARC1001339	32.829	42.256	31.603	10.158	27. 332	21,573	35. 452	25. 220
	OYARC1001340	27.630	18.361	12.822	7.427	6.739	12.500	23, 923	14. 457
	QYARC1001341	95, 252	81.979	52.630	68. 282	53.071	55, 813	59.589	60.054
	***************************************		252.091		202. 538	60.427	87. 325	80. 221	137.940
	OYARC1001342	100.966						56.015	68. 157
	OYARC1001344	103.513	107.791	75. 126	75.888	55. 791	47. 394		
	OVARC1001357	10.771	20.444	6.064	5. 959	2.545	8. 202	6.654	9. 212
	OVARC1001359	74.406	41.612	39.409	39. 521	22.602	47.817	49, 919	41.248
	OYARC1001360	12.963	15.729	5. 865	8. 162	5. 343	8.344	7.449	5. 231
	OVARC1001369	30.741	30.024	17. 593	14. 376	15.376	19.395	28. 970	17, 236
				28. 420	22.363	23, 533	27.224	35.738	26. 351
	OVARC1001372	47.372	31.878			52.979	38.758	43. 990	55.762
	OVARC1001376	65.628	113.295	43.890	78.146				
	OVARC1001381	115.063	118.072	70.088	92. 127	69.013	60.845	53.880	62.779
	OYARC1001391	39.498	37.024	30.883	12.771	21.036	26.802	26.851	18. 964
	OVARC1001392	17.841	15.639	29.498	12.487	18.354	13, 407	13.843	15. 944
	OYARC1001399	43, 831	87,706	37.282	44. 533	34.853	26, 357	28.943	38.749
	OYARC1001417		24.005	20.041	15. 997	12, 488	15. 218	23.379	13. 202
		26.403		47.763	31.720	43, 415	60.531	66.782	40.700
	OYARC1001419	102.361	46.760			18.684	27. 353	39.805	22.560
	OYARC1001425	36.511	32.857	19.181	27.837				
	OVARC1001436	56.321	33. 132	21.728	19,600	24.952	23.512	43. 382	21. 101
	OVARC1001442	85.715	36.595	24.645	21.266	30.507	37.805	58, 999	27, 499
	OVARC1001451	34.303	30.697	30.804	34.477	24. 521	23.798		24. 423
	OYARC1001452	53, 317	30.445	17. 186	12.444	18.765	27. 539	29.572	22. 163
	OYARCIGO1453	16.620	33. 383	8.673	8.363	7.911	7.294	15.113	10.726
	OVARC1001476	23, 408	34.645	23.709	17. 349	17.688	17.078	15, 241	27. 167
					21.037	14.968	36. 453	52.487	28. 092
	OVARC1001480	69.410	32. 323	28. 385			4.046	10, 274	9. 908
	OVARC1001489	10.998	9. 249	5. 028	7.129	6. 338			
	DVARC1001493	55. 166	55. 346	14. 849	9, 601	15, 915	27.767	38.065	22. 112
	OVARC1001496	85.220	65.108	29.250	24, 050		36, 194	61.219	38. 523
	OVARC1001499	27.560	27.910	16.669	16.239		20.989	35. 173	22. 472
	OVARC1001506	67.326		43, 800	30.337	35.006	34, 184	46.403	31. 327
				29.884			23, 100		18.605
	OVARC1001509	45, 793				11.932	6.535		9. 275
	OVARC1001510	14.065		12.458					
	OVARC1001516	64. 781	44. 167	26.084			33.509	55. 926	34.068
	OVARC1001525	8, 675	10.658	6.559	6.011		3. 863		5. 884
	OVARC1001542	34.447					20.034	32.156	28. 167
							35, 307		51.833
	OYARC1001544	97.739							34.075
	OVARC1001546	40.692							
	OVARC1001547	4.108							5. 293
	OVARC1001555	48.644	45.769	33.072	18. 823	21.553	30, 921	53.633	33.664
		1							

Table 124

	INCLOSION I	0.006	11.616	22.248	5.899	8. 179	16.185	18, 151	7.957
	OVARC1001560	9.995			17.742	13.831	32.806	32, 162	24.321
	OVARC1001569	40.746	31. 448	15.414				50. 455	32.827
5	OVARC1001570	45.828	32. 466	28.804	16.797	17.223	29. 282		
•	OVARC1001577	18.703	19.196	13.453	17. 108	9.651	14.718	23.685	24.544
	OVARC1001578	4.894	1.347	3. 487	1.668	2.647	3.022	0.000	0.000
	OVARC1001596	84.296	49.737	31.737	18.041	23.005	61. 151	47.274	34.947
	OVARC1001600	54.416	43. 232	24.561	22.726	16.594	23.734	27.443	19.377
	OVARC1001607	21.077	19.469	15.218	12.687	5.720	14, 273	22. 223	16.907
			13.445	7.606	4.839	6.723	8. 590	14.535	14.413
10	OVARC1001610	22.320		11.190	3.816	6. 271	10. 248	18. 405	10. 394
	OVARC1001611	10.788	15. 290				39. 578	60. 842	26. 422
	OVARC1001615	83.171	33.856	33, 256	23.489	27. 385			
	OVARC1001636	19. 126	18. 265	9, 929	10.903	5.896	14. 319	20.083	11. 921
	OYARC1001658	184.639	178. 409		130.922	77.578	71.883	78.800	99.902
	OVARC1001702	74.853	43. 682	37.735	17. 471	24. 833	47.858	40.347	30.531
	OVARC1001703	20.271	16.866	19.593	10.314	12. 106	14. 193	17. 305	11.237
15	OVARC1001710	104.705	53. 627	45.081	22. 841	30.909	61.922	\$7.754	30.671
	OVARC1001711	38.919	48. 731	30.797	20.615	17. 927	29.742	29.051	30. 493
	OVARC1001713	58, 871	50.075	38.715	24.728	28.026	37.714	59. 338	46.137
	OVARC1001725	12.462	6. 462,	9.161	5.766	5. 579	7.643	12. 283	11.952
	OVARC1001726	60. 846	30. 421	22.951	16. 102	17, 141	25. 341	40.000	23.764
		12.749	6. 695	1.629	3. 384	2.943	5. 347	11.864	5.882
20	OVARC1001727		296. 389	159, 879	90.412	79.927	104. 739	112.601	182.645
	OVARC1001731	417.237		13.004	10.850	7779	19. 246	25. 926	9.776
	OVARC1001735	29. 333	21.981		40.754			40. 353	35.965
	OVARC1001741	62. 439	80.254	36. 924		30. 175	31. 693 42. 496	52.805	40.912
	OVARC1001745	105. 943	90. 392	54.073	48. 385	29. 915			
	OVARC1001759	6. 344	6, 101	7, 549	5.672	5. 285	7.629	4. 284	16.699
	OVARC1001762	15.752	20.242	8.966	13. 129	12. 132	11. 198	17.879	12.812
25	OVARC1001766	50. 421	44.814	32.524	34.415	32.044	28. 483	25.974	25. 134
	OVARC1001767	12.694	11.424	7.232	4. 392	4. 561	7.783	6.753	3.775
	OVARC1001768	30.851	32.866	18.111	12. 623	14.716	15.800	18. 499	17.641
	OVARC1001770	99.967	29.814	24.915	16.646	18. 553	49.766	33.065	24.957
	OVARC1001776	84.733	66.614	35, 351	18.038	20. 855	40. 357	40.259	38.765
	OVARC1001791	82.228	59. 107	38.878	27.000	19.647	41.210	58. 352	34.508
30	OVARC1001795	35, 170	31.032	19,091	14.053	19.096	16.818	23.677	24,540
55	OVARC1001798	113.936	95.099	73, 266	84.613	71.384	61.440	58. 197	68, 577
		125.877	98.941	72.747	75. 225	59. 196	77.683	67.227	71.441
	OVARC1001802		10.835	12.686	8.980	8. 339	13.601	7.696	8.902
	OVARC1001805	10. 464	172.138	42, 410	25. 456	42. 245	77.908	59.683	39, 476
	OYARC1001807	135.513				49.795	56.085	64.919	59.018
	OVARC1001809	118. 235	105. 836	62.430	46.885		38. 322	38.347	31.540
35	OVARC1001812	67. 287	48.010	53.706	41.376	36. 383			36.514
	OVARC1001813	69.943	84, 621	53.953	55. 458	42.844	41.002	32.364	
	OYARC1001820	52.381	53.833	35.503	41.319	24.742	28.840	25.646	28.845
	OYARC1001828	8.200	10.217	4.364	9.812	6. 280	8.885	6.886	8.407
	OVARC1001833	86.833	60.894	37.693	22.705	29.730	50. 489	52.516	40.092
	OVARC1001839	39.140	38, 162	14. 245	19.805	17.227	23. 521	26.722	22.628
40	OVARC1001846	14.794	24.500	15, 503	10.407	8. 977	15. 603	9.900	14.219
70	OVARC1001849	73.011	60.883	43.536	39.792	33.900	30. 397	28. 153	30.952
	OVARC1001861	61.938	43.449	26.931	16.558	17. 111	24.800	36.196	21.959
	OVARC1001873	37.219	38.842	19.844	22. 293	20.314	24. 148	34. 160	26.819
	OVARC1001879	76.088	51.361	39.655	29.363	28.800	45.644	47.894	29.618
	OYARC1001880	135, 860	84.254	58.296	66,680	55. 691	73.306	83.823	57, 413
	OVARC1001883	81.852	74. 425	52.983	53.494	55. 481	39,665	45.082	50.587
45	OVARC1001900	55. 149	42.744	20.659	17. 501	28.891	25. 216	36.722	27.567
	OVARC1001901	35. 402	43, 250	19.139	18.068	14.966	16, 860	28.327	21.865
				16.048	15.000	9. 189	16.480	16. 595	14.072
	OVARC1001911	26.676	31.540			24. 346	38. 467	49,017	28.751
	OVARC1001916	57.008	57.583	30.437	33. 497			7. 928	
	OYARC1001928	11.760	11.451	9.871	8. 924	3. 218	9.310	28.632	8.861 31.167
50	OVARC1001937	41.094	331.797	26. 182		18.612	29. 201		
00	OVARC1001940		25.633	19.059		15. 166	24. 914	25. 701	31.361
	OVARC1001942	30.967	37. 334	26.741	17.951	21.439	17. 640	29.921	25.107
	OVARC1001943	85. 434	52. 979			35.086	45. 562	49.703	36.562
	OVARC1001949	27.732		29.233		22.996	24.826	26. 681	54.991
	OVARC1001950	114.630	90.867			43. 996	70.058	71.925	46.593
	OVARC1001952	140.095		76.000	76.812	\$7.544	76, 600	127. 024	117. 497
55									

Table 125

		-0.10	34 364 1	24.826	18,570	18.070	29. 136	28.466	23.093
	OVARC1001954	38. 148	34. 154		43, 880	39. 594	41.763	47.995	42.856
	OVARC1001963	70.685	73.510	38. 247	104. 971	53. 259	65. 630	60,001	74.940
_	OVARC1001983	103.819	85.974			28. 833	28.724	72.176	33, 793
5	OVARC1001987	55.904	47.294	16.298	23. 921	109.717	72.479	80. 807	76.628
	OVARC1001989		123.408		101.800				50, 807
	OVARC1001991	105.789	61.566	52.852	26.772	46.555	61.290	57. 420	
	OVARC1002005	43.909	50, 446	52.235	34, 217	36.792	22. 115	37. 361	38. 275
	0YARC1002044	68. 989	92.088	47.242	50.982	37.959	40. 246	32.518	39. 591
	OVARC1002046	142.697	103.646	61.978	48.709	50.959	89.078	107.957	65. 922
)	OVARC1002050	150.418	79.832	52. 259	30.717	50.113	74.307	95. 763	52.005
	OVARC1002058	31.725	28, 126	16.056	14. 204	10. 314	24.705	25.418	21.662
	OVARC1002066	22.845	30.065	5. 783	10.572	14.029	17. 339	17.127	29, 452
	OVARC1002082	142.891	132.300	69.068	83.947	73.662	57.050	101.160	85. 803
	0VARC1002091	49. 223	45.691	28. 357	26.618	29. 431	30.906	42.116	35. 122
	OVARC1002092	16. 502	13.059	11.455	8.985	10.442	10.939	11.841	11.658
		206, 510	229. 583	94, 978	52.679	65.398	105.804	108.886	55.303
	OVARC1002093		70.842	22, 175	21.837	26. 392	25. 855	44.104	28. 562
	OVARC1002094	57. 983		51.719	88.540	57.860	40.826	46.289	44,621
	DVARC1002107	81. 163	81.383		85. 579	47. 248	39.907	60.603	67, 156
	OVARC1002112	71. 336	80.431,	40. 320	35.010	40. 692	63.760	106.294	65. 520
	OVARC1002126	114. 239	87.851	47.175		21.021	26. 217	51.525	32. 857
	OYARC1002127	55. 311	43.006	22.728	10.831	8. 159	7. 853	10.350	10, 188
	OVARC1002138	8. 951	13.827	3.935	7.856			31.142	20, 410
	OVARC1002143	46.546	34.713	16.566	15. 769	19. 276 8. 291	24.331 12.374	13.614	15.810
	OVARC1002156	12.544	23.040	10.035	15.363			31. 215	19.064
	OVARC1002158	56. 221	28.255	18. 260	10.748	16. 251	20.791		75, 063
	OVARC1002165	101. 989	143.172	84.011	82.086	75. 946	58.837	85. 203	
	OVARC1002175	207. 395	83.881	84.413	54.135	104.278	114.458	134. 235	69. 297
	DVARC1002178	17.313	27.443	12.750	10.705	15. 530	12.936	23.362	17.872
	OVARC1002182	40. 283	37.762	18.779	11.770	18.311	21.416	34.402	24. 309
	OVARC1002185	36.278	33.563	17. 925	17. 394	20.095	28. 241	45. 498	31.989
	PLACE 1000004	41.829	37.799	18. 473	16.218	12.661	20. 372	25,010	22.000
	PLACE 1000005	33, 315	36.712	26.079	24.859	17.404	25.038	28. 162	24.028
	PLACE 1000008	48. 081	38.647	24. 284	19.081	18.255	32.116	54, 951	30. 255
	PLACE 1000007	24. 221	25.983	17.339	11.998	16. 921	17.706	46.581	21.338
•	PLACE 1000014	57.292	49.432	36, 234	32.812	25. 276	24.815	35. 655	32.759
	PLACE1000031	42.309	61.878	50. 107	46.094	37.373	29.757	38. 437	47.194
	PLACE 1000033	7.856	22.257	8.411	7.606	9.169	11.609	12.768	10.286
	PLACE 1000040	36.717	30.479	20. 358	21.457	23.948	12.296	22.459	20.099
	PLACE 1000048	32. 105	28. 302	21.619	18.209	13.458	16.364	16.026	12.244
	PLACE 1000050	33. 955	41.358	21.915	18. 172	15. 208	24.691	30.515	22.038
ī	PLACE 1000061	159. 492	228.723	82.722	177.569	132, 119	143.553	116. 181	188, 103
	PLACE 1000066	59. 266	55.710	42.829	38.851	45.700	47, 171	50. 185	56.938
	PLACE 1000075	15. 690	15.994	12.949	6.500	11.914	10.574	6.929	11.391
	PLACE 1000078	46. 952	57.637	52.225	42.480	22.126	28.527	38.463	41.033
			63. 282		23.924		30. 920	50.546	41.886
	N. 1961VVVVV					1 23.114	1 30.320		
	PLACE1000081	75. 884		38.644 52.586		29.174 38.070	64.488	55. 431	42.640
)	PLACE1000086	85. 184	67.162	52.586	27.421	38.070	64.488	55. 431	
,	PLACE1000086 PLACE1000094	85. 184 49. 828	67.162 42.276	52. 586 20. 226	27.421 10.189	38.070 9.355	64.488 21.041	55. 431 19. 625	42.640 8.506
)	PLACE1000086 PLACE1000094 PLACE1000101	85. 184 49. 828 10. 188	67. 162 42. 276 23. 449	52. 586 20. 226 16. 699	27. 421 10. 189 19. 362	38.070 9.355 17.073	64.488 21.041 11.091	55. 431 19. 625 13. 623	42.640
)	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000121	85. 184 49. 828 10. 188 56. 678	67. 162 42. 276 23. 449 34. 412	52. 586 20. 226 16. 699 30. 070	27. 421 10. 189 19. 362 13. 506	38.070 9.355 17.073 19.044	64.488 21.041 11.091 31.104	55. 431 19. 625 13. 623 40. 290	42.640 8.506 19.675
0	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000121 PLACE1000133	85. 184 49. 828 10. 188 56. 678 39. 057	67. 162 42. 276 23. 449 34. 412 29. 915	52. 586 20. 226 16. 699 30. 070 23. 128	27. 421 10. 189 19. 362 13. 506 29. 843	38.070 9.355 17.073 19.044 20.718	64.488 21.041 11.091 31.104 24.672	55. 431 19. 625 13. 623 40. 290 26. 803	42.640 8.506 19.675 25.078 39.107
0	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000121 PLACE1000133 PLACE1000142	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740	38.070 9.355 17.073 19.044 20.718 21.065	64. 488 21.041 11.091 31.104 24.672 43.454	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693	42.640 8.506 19.675 25.078 39.107 35.205
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834	67.162 42.276 23.449 34.412 29.915 47.628 62.270	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131	38.070 9.355 17.073 19.044 20.718 21.065 27.243	64.488 21.041 11.091 31.104 24.672 43.454 17.729	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001	42.640 8.506 19.675 25.078 39.107 35.205 47.833
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000121 PLACE1000132 PLACE1000142 PLACE1000146 PLACE1000163	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015	67.162 42.276 23.449 34.412 29.915 47.628 62.270 87.206	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.941	64.488 21.041 11.091 31.104 24.672 43.454 17.729 62.968	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001 120. 625	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000121 PLACE1000133 PLACE1000146 PLACE1000163 PLACE1000172	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508	67.162 42.276 23.449 34.412 29.915 47.628 62.270 87.206 23.847	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595	38.070 9.355 17.073 19.044 20.718 21.085 27.243 45.943 10.491	64.488 21.041 11.091 31.104 24.672 43.454 17.729 62.968 12.594	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001 120. 625 4. 745	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000163 PLACE1000176 PLACE1000181	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850	64.488 21.041 11.091 31.104 24.672 43.454 17.729 62.968 12.594 23.966	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001 120. 625 4. 745 21. 392	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000121 PLACE1000133 PLACE1000146 PLACE1000163 PLACE1000172	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 6. 68	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 4 2. 195	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764	64.488 21.041 11.091 31.104 24.672 43.454 17.729 62.968 12.594 23.966 24.786	55. 431 19. 625 13. 623 40. 290 26. 803 39. 001 120. 625 4. 745 21. 392 5. 246	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000163 PLACE1000176 PLACE1000181	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 6. 684 41. 261	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 4 2. 195 26. 145	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 12. 594 23. 966 24. 786 35. 082	55. 431 19. 625 13. 623 40. 290 26. 803 39. 001 120. 625 4. 745 21. 392 5. 246	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199
	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000163 PLACE1000181 PLACE1000181	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226 45. 178	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 41. 261	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 42. 195 26. 145	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092 14.083	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 12. 594 23. 966 24. 786 35. 082	55. 431 19. 625 13. 623 40. 290 26. 803 51. 693 39. 001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227
5	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000172 PLACE1000181 PLACE1000185 PLACE1000185	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981 34. 090	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226 45. 178 28. 795	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 41. 261	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 4 2. 195 26. 145 10. 196	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092 14.083 20.362	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 12. 594 23. 966 24. 786 35. 082 15. 181 61. 122	55. 431 19. 625 13. 623 40. 290 26. 803 51. 693 39. 001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504 55. 368	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227 17.891
5	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000163 PLACE1000172 PLACE1000181 PLACE1000181 PLACE1000185 PLACE1000186 PLACE10001898 PLACE1000198	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981 34. 090 29. 427	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226 45. 178 28. 795	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 41. 261 19. 770 20. 16	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 4 2. 195 26. 145 0 10. 196 17. 037	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092 14.083 20.362	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 12. 594 23. 966 24. 786 35. 082 15. 181 61. 122	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39.001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504 55. 368	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227 17.891 2.408
5	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000172 PLACE1000181 PLACE1000184 PLACE1000185 PLACE1000181 PLACE1000181 PLACE1000181 PLACE1000181 PLACE1000181	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981 34. 090 29. 427 8. 728	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226 45. 178 28. 795 38. 826	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 6. 584 41. 26 19. 770 20. 161 9. 733	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 12. 195 26. 145 0 10. 196 17. 037 11. 456	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092 14.083 20.362 12.426	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 12. 594 23. 966 24. 786 35. 082 15. 181 61. 122 6. 184	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504 8. 011 26. 274	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227 17.891 2.408 20.792
10 15	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000172 PLACE1000181 PLACE1000184 PLACE1000185 PLACE1000188 PLACE1000181 PLACE1000181 PLACE1000181 PLACE1000181 PLACE1000181	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981 34. 090 8. 728 35. 035	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 23. 847 36. 469 3. 226 45. 178 28. 795 38. 826 14. 768	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 6. 684 41. 261 19. 770 20. 161 9. 731 22. 38	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 1 2. 195 26. 145 0 10. 196 17. 037 3 11. 456 7 16. 421	38.070 9.355 17.073 19.044 20.718 21.065 27.243 10.491 30.850 8.764 25.092 14.083 20.362 12.426 20.597	64. 488 21.041 11.091 31.104 24.672 43. 454 17.729 62.968 12.594 23.966 24.786 35.082 15.181 61.122 6.184 25.167	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504 8. 011 26. 274	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227 17.891 2.408 20.792
15	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000172 PLACE1000181 PLACE1000184 PLACE1000185 PLACE1000188 PLACE1000198 PLACE1000213 PLACE1000213 PLACE1000220 PLACE1000220	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981 34. 090 29. 427 8. 728 35. 035 348. 135	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226 45. 178 28. 795 14. 768 36. 902 182. 545	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 41. 26 19. 770 20. 161 9. 73 22. 38 114. 75	27. 421 10. 189 19. 362 13. 506 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 1 2. 195 1 26. 145 0 10. 196 7 16. 421 5 86. 687	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092 14.083 20.362 20.597	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 22. 966 24. 786 35. 082 15. 181 61. 122 6. 184 25. 167	55. 431 19. 625 13. 623 40. 290 26. 803 61. 693 39. 001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504 8. 011 26. 274 106. 589	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227 17.891 2.408 20.792 98.294
15	PLACE1000086 PLACE1000094 PLACE1000101 PLACE1000133 PLACE1000142 PLACE1000146 PLACE1000172 PLACE1000181 PLACE1000184 PLACE1000185 PLACE1000188 PLACE1000181 PLACE1000181 PLACE1000181 PLACE1000181 PLACE1000181	85. 184 49. 828 10. 188 56. 678 39. 057 59. 811 68. 834 102. 015 9. 508 51. 412 16. 961 62. 981 34. 090 29. 427 8. 728 35. 035 348. 135 79. 604	67. 162 42. 276 23. 449 34. 412 29. 915 47. 628 62. 270 87. 206 23. 847 36. 469 3. 226 45. 178 28. 795 38. 826 14. 768 36. 902 182. 545 63. 001	52. 586 20. 226 16. 699 30. 070 23. 128 31. 984 34. 321 45. 923 6. 470 31. 628 41. 261 19. 770 20. 161 9. 73: 22. 38: 114. 75: 31. 91	27. 421 10. 189 19. 362 29. 843 14. 740 47. 131 38. 164 9. 595 23. 060 4 2. 195 10. 196 17. 037 31. 1. 456 7 16. 421 5 86. 687 9 29. 088	38.070 9.355 17.073 19.044 20.718 21.065 27.243 45.943 10.491 30.850 8.764 25.092 14.083 20.362 12.426 20.597 95.201	64. 488 21. 041 11. 091 31. 104 24. 672 43. 454 17. 729 62. 968 12. 594 23. 966 24. 786 35. 082 15. 181 61. 122 6. 184 25. 167 164. 292	55. 431 19. 625 13. 623 40. 290 26. 803 39. 001 120. 625 4. 745 21. 392 5. 246 37. 231 22. 504 55. 368 8. 011 26. 274 106. 589 26. 593	42.640 8.506 19.675 25.078 39.107 35.205 47.833 52.326 20.625 24.437 4.794 52.199 21.227 17.891 2.408 20.792 98.294 31.426

Table 126 .

	C	62 620 I	60,061	23.370	26.968	16, 702	28.102	27.116	50.991
	PLACE 1000248	63.620	86.542		92.906	67.210	62.207	74.824	84. 158
		07. 386			51. 291	33.764	52.872		56.824
_	PLACE 1000288		30.859		76. 783	64. 315	53.082	45.786	64.840
5	I Diocionage		07.978	64.652		73. 167	20.044	16.870	23.303
	PLACE 1000302	36.212	31.35	48. 891	8. 192	21.314	32.791	44. 370	34, 969
	PLACE1000304	77.695	50.861	24.615	19.705		11.170	8, 490	10.525
	PLACE1000308	13.844	18.591	10.915	15. 228	13.497		89.605	51. 438
	PLACE1000309	171.086	79.282	53.477	30.661	44. 221	99.582	12.364	24.742
	PLACE1000312	25.013	29.701	14.081	15. 125	7.699	11.121	19. 395	12.431
10	PLACE1000330	29.657	13. 102	12.306	10. 127	9. 659	16.951		8. 302
	PLACE1000332	13. 294	6.752	7.366	6. 938	4. 823	5. 141	7.821	
	PLACE 1000347	45.531	37. 378	19.406	17. 234	19. 477	19.786	29. 460	24, 427
	PLACE 1000351	93. 299	56.437	40.461	27. 466	26. 428	44.784	56. 685	47.749
	PLACE 1000374	89.871	66.668	53. 557	66,616	45.909	45.689	49.979	76. 296
	PLACE 1000380	22.012	21.037	15.351	9. 985	12.229	7.428	19.713	17.050
15	PLACE 1000383	29.005	24.752	16.349	15. 183	11.959	16.827	29. 293	19.713
	PLACE 1000397	35. 368	26. 208	19.042	6, 636	9.008	19.143	19.667	12.826
	PLACE 1000401	121.012	77.115	91.986	73.017	85. 204	77.208	97.740	89.014
	PLACE 1000406	43. 944	37.883	20.305	20. 530	17.412	26,601	31.601	28. 177
	PLACE 1000412	14.283	24.094	8. 322	10. 240	8. 659	9.837	13.298	15. 409
	PLACE 1000420	95. 364	99. 949	57. 598	50. 129	39. 257	39. 215	34.611	48. 196
20	PLACE 1000421	59.754	60. 388	52.953	37. 350	31.433	40.619	40.095	47.679
20	PLACE1000423	49. 130	51.837	22.800	9.952	21.218	55, 558	49.895	90. 332
	PLACE 1000424	57. 584	46. 928	23.243	25. 445	11.122	23. 277	21.409	24. 420
	PLACE 1000430	9,008	14. 497	9.324	6.648	12.506	3. 176	4. 529	9.751
	PLACE 1000433	61.817	33.422	22.755	10.220	12.877	30.460	38.040	20.834
	PLACE 1000435	58. 251	49.033	24, 534	33. 925	25. 131	18.878	26.453	33.894
05	PLACE 1000437	37.691	42.505	19, 354	13.578	22. 936	27.017	52.788	24.766
25	PLACE 1000442	28.959	53.891	36.443	45.883	36.218	29.092	28.915	59.563
	PLACE 1000444	222.629	268, 192	164.724	178.057	143.884	113, 247	125.051	157.345
	PLACE 1000453	60, 912	56.717	45. 737	28.913	34.374	46. 491	47.877	46.094
	PLACE 1000456	59.850	55.649	26, 148	10.788	16.900	32.811	31.014	25. 102
	PLACE 1000465	63.781	32.184	28.609	22.813	15.851	25. 834	76.172	29.680
	PLACE 1000481	117.442	55.048	43.008	40.607	39.135	57.771	52.403	44. 241
30	PLACE 1000492	85. 199	42.804	28.200	13.820	16. 493	35.818	62.470	37.299
	PLACE 1000508	48. 116	30.697	17.662	19. 193	14. 645	26. 357	39.846	23.454
	PLACE 1000512	23.066	37.331	52.438	15.899	43.633	17. 392	16.605	25.441
	PLACE 1000540	6. 354	22.237	6.827	9.533	6.338	8. 582	5.690	8.570
	PLACE 1000541	139. 592	95.891	62.856	44. 350	48.779	102.808	118.737	83.454
	PLACE 1000546	24.434	15.843	9.613	13.003	8. 921	13.653	21.807	14.697
35	PLACE 1000547	138.587	72.254	64.656	57.672	49.694	71.928	84.849	56.997
	PLACE 1000560	39.727	25.726	19.961	10.708	14, 907	22. 472	42.419	29.563
	PLACE 1000562	74.380	77.139	35.608	44.686	31. 444	29.868	26.773	50.026
	PLACE 1000564	45.712	39.050	20. 165	14.663	19. 526	22.670	43.140	35. 028
	PLACE 1000583	122.345	132.820	73.526	90. 516	75. 343	62.557	52. 925	95.075
	PLACE 1000587	99.842	63.364	42.075	55. 988	38.170	36.599	30.062	35.245
40	PLACE 1000588	86, 166	135.917	34.894	41.374	26.506	42.479	60.642	72.805
	PLACE 1000596	49. 265	55. 996	23.832	26.469	29.318	57.681	28.073	35.812
	PLACE1000599	79, 259	72.325	37.975	49.064	36.704	32, 501	40.446	38. 539
	PLACE 1000605	46.938	54.185	20.654	19.011	15. 275	25. 549	73.210	37.742
	PLACE 10006 10	45.555	31, 108	16.017	11.318	13.984	22.493	36.775	27.839
	PLACE 1000611	83.806	12.237	34.984	19.496	31.956	36.823	73.743	37.315
45	PLACE 1000626	25, 444	20.294	26.796	13.307				24.644
	PLACE1000633	51.819			36.919	35. 957			40.217
	PLACE 1000636	19.979		10.22	15.590	13.380			13.996
	PLACE 1000653	19, 174				12.647			11.318
	PLACE 1000656	207.889					78.748		
	PLACE 1800663	27. 908					14. 462	23.389	
50	PLACE 1000706	283.571						210.569	73.830
		61,631						57.877	39.148
	PLACE 1000712								15.949
	PLACE 1000716	26.011							
	PLACE 1000740	34, 490							
	PLACE1000748								
55	PLACE 1000749	246.155	130.04	101.03	J 10.31	. , , , , , ,			

Table 127

	(D) AAC (AAAYE)	8.591	28.632	6.888	6.628	8.859	7.678	7.925	11.115
	PLACE 1000751				14. 166	9, 125	11.761	20.466	14. 904
	PLACE 1000755	22.080	22.789			5. 547	12.731	21.034	18.074
5	PLACE 1000769	16.024	19. 119		12. 207				
3	PLACE1000778	109.940	168.867	46.116	36, 217	54. 573	33. 450	40.021	61.410
	PLACE 1000785	54.501	35, 590		21.344	11.939	17. 233	19.818	30.628
	PLACE1000786	63.401	34.818	26.260	25.783	21. 236	33. 236	29.738	24.419
	PLACE 1000793	48.092	49.470	31.204	14. 276	17.894	36.450	38.082	31.337
	PLACE 1000795	38. 178	43, 688	29.889	21.674	10.765	21.955	41.921	41.550
	PLACE 1000798	31.236	40, 770	22.606	25. 191	17. 921	17, 856	21.782	21.758
10			23. 549	17. 121	14.965	8.140	11.726	13.094	25.608
	PLACE 1000812	24. 169		40.416	88, 702	37. 795	36.623	32.882	44.655
	PLACE 1000823	81.457	78.801			31.956	47.353	61.449	155.007
	PLACE 1000825	72.220	107. 715		34. 346			59. 936	33.016
	PLACE 1000838	44.642	81.659	25.304	15. 146	16.808	62.951		
	PLACE 1000841	19.731	9. 168	3. 325	14.206	7.817	19.073	12.783	10.985
	PLACE 1000843	23.326	27.970	19.816	10.746	10.401	15. 372	21.004	17. 198
15	PLACE 1000849	171.333	62.539	65.353	38.857	44. 284	93.732	118.022	61.526
	PLACE 1000856	35.302	20. 267	19.938	11.857	16.559	18. 275	24.314	14.524
	PLACE 1000863	61.947	24.729	17.318	10.548	32.356	45.071	40.695	26.491
	PLACE 1000876	79.589	41.303	31.803	27, 682	30.566	41.161	58.457	35.801
	PLACE 1000899	36.028	54. 514	19.200	14. 563	23.816	17, 191	23.052	18.916
	PLACE 1000893	34.468	58. 737	45.762	53. 355	33.953	45, 837	23.987	112.516
20		17.260	18. 289	7.853	7,770	4.100	9. 541	11.860	5. 411
	PLACE 1000909				18. 579	21.304	47.829	47.423	31.816
	PLACE 1000912	72.300	41.738	29.873			13,771	20. 247	22.212
	PLACE 1000914	34. 274	20.778	16.170	8.631	12.137			7.614
	PLACE1000918	6.646	24. 953	6.298	21.039	6.076	8.001	14. 538	
	PLACE1000927	28.004	62.278	11.519	25. 240	15. 288	27. 303	24.639	36.302
	PLACE 1000931	60.013	70.374	41.114	48. 090°	31.983	37.593	35.750	38. 045
25	PLACE1000944	15.469	20.100	11.329	9. 563	11.301	10.496	10.907	13.479
	PLACE1000948	32.119	174.384	19.147	11.561	8.864	13.716	16.344	23.012
	PLACE 1000958	24.559	28.912	20, 683	12, 101	15.980	17.758	18.551	16. 227
	PLACE 1000 972	120.934	92. 540	56.098	49, 344	42.043	64.933	66.648	83.486
	PLACE 1000977	5.160	23.071	5. 930	7, 450	7.265	6.795	7.348	13.664
	PLACE 1000 979	36.518	36.872	31.314	43.863	34.967	34.693	38.011	58, 543
30	PLACE 1000 986	39.462	32, 248	17.759	9. 962	10.922	17.210	20.134	11.670
		85.543	56.030	33.710	26.097	53.247	35, 833	43.907	42.264
	PLACE 1000 987			11.199	15, 991	10.697	10.336	15.117	15.657
	PLACE 100 1000	15.969	18. 182		21.556	18.037	24, 959	24.887	38.857
	PLACE 100 1007	41.857	48. 683	23.082		14.359	16.726	17.763	18. 489
	PLACE 1001010	29.468	27. 943	26. 350	21.964		13.946	8.904	23. 581
25	PLACE 1001015	20.540	30,643	16.387	20.211	7.569	36.963	57.898	62.858
35	PLACE 1001016	77.787	52.441	29, 862	30. 282	22.094			20.499
	PLACE 1001022	33. 101	30. 827	19. 383	11, 100	14.872	22.156	23.755	
	PLACE 100 1024	86.274	27.421	25, 662	12.087	19.171	38. 266	41.922	20.735
	PLACE 100 1036	80.642	165. 022	64. 983	46.681	67.747	121.474	402. 289	252.956
	PLACE 1001038	452.345	139. 825	89.101	67.372	64. 392	122.656	119.479	107.665
	PLACE 1001048	49.948	49.581	16.660	14. 592	10.687	24.644	16.889	36. 435
40	PLACE1001054	134.306	67.365	61.474	36.835	33.520	69, 944	111.570	67.974
	PLACE 1001062	74.158	68.783	52. 589	64.589	49.941	41.816	51.497	54.685
	PLACE 1001063	10.880	13.653	8.862	9.859	6.427	6.510	8.010	9.447
	PLACE 1001076	14.575	15.670	12.223	5.950	12.881	9.910	15, 204	12.067
	PLACE 1001081	12. 530	13. 285	8.314	5.016	4.852	19.472	31.441	10.426
	PLACE 100 1088	25.759	16. 332	10.811	9. 362	11.626	15. 207	22.359	14.210
45	PLACE 100 1092	15, 938	44. 121	18. 940	15, 854	15.358	18.646	27.718	25.006
45		51.863		44. 477	36.802	35.002	36,534	40.789	44.072
	PLACE 100 1098		74. 664			21.199	38.215	39.752	36.621
	PLACE 100 1 100	69.984	61.458	42.513	19, 257	15.200	22, 158	23.976	24.947
	PLACE 1001 104	37.879	43.589	22.459			22.370	24.583	26.608
	PLACE 1001114	50.995	43. 129	28.583	41.340	23.689			
	PLACE 1001118	55.858	39, 536	30.416	29. 284	13.566	35.583	35.042	61.564
50	PLACE 100 1 123	30.236	32.692	12.932	16.066	9.901	19. 213	20.910	28.778
	PLACE 100 1 136	127. 205	106.279	47.874	46.520	45. 126	54.639	67.043	60.071
	PLACE 1001144	59. 577	14.773	33.377	21.823	38.443	32.412	40.190	39.315
	PLACE 1001147	59.813	42.869	27.085	20.092	30. 181	39. 398	45, 339	34.463
	PLACE 1001148	37.059	29.368	18. 220	13. 240	14.014	19.609		28.919
	PLACE 100 1159	23.780	18.761	10.274	9. 929	12.302	17. 285	19.282	17.753
	PLACE 1001168	26.768	24. 323	12.289	8.468	8.558	14.711	22.168	20.921
55									

Table 128

				-A -116 T	0 700 1	13 646 1	20. 994	34.674	26. 387
	PLACE 1001171	37.609	26.312	19.416	9.788	11.645			
	PLACE1001183	48.472	34. 255	16. 988	12. 402	13.998	24.043	41.590	34_738
	PLACE1001185	98, 156	72.026	33.520	17. 455	34.874	41. 246	78. 451	45, 433
5			28. 202	14.832	19. 137	15.156	11.504	21.093	18.878
•	PLACE1001201	20.710			24.039	19.810	29.842	30.368	33.000
	PLACE1001229	33. 202	50.727					30.611	22. 184
	PLACE 1001231	28.893	32.022	21.470	16.244	15.489	23.482		
	PLACE 1001238	67.072	60.114	37. 423	43. 278	30.120	34.706	41.011	38. 313
	PLACE 1001241	21,610	25.407	7. 984	17. 578	8.443	14.781	31.035	14. 575
			69.441	28. 266	26.878	24.774	29. 386	68. 093	55. 636
10	PLACE 1001242	45, 592		7. 186	6.906	8.128	9.488	6.808	15. 989
,,,	PLACE 1001247	14. 525	18. 387			20.092	23.448	40. 190	18. 900
	PLACE 1001250	49.114	30.049	15. 521	12. 388			37. 267	51. 354
	PLACE1001257	62.294	83.027	38.705	44. 550	45.672	38. 236		
	PLACE1001272	63. 255	35.776	22.716	18.567	23.479	28.934	54. 496	33. 742
		20.477	21.478	8. 935	8. 448	12.817	12.013	16. 223	11. 151_}
	PLACE 1001279		56.354	46.699	32.609	50.557	37.478	30.514	34. 496
15	PLACE1001280	68.512				22.103	14. 441	14. 208	23. 363
13	PLACE1001294	16.622	36. 599	12.414	23. 498			110. 093	39. 852
	PLACE 1001295	158.866	63.791	43.310	26.850	56.659	72.706		
	PLACE 1001300	64, 491	33, 466	14.714	9. 167	18.136	13.210	28. 528	23. 798
	PLACE1001304	70.999	60.035,	54.352	72.559	49.765	40.745	55.843	97. 914
		77.711	67.514	37.479	36.657	50.824	35. 191	38. 273	47. 028
	PLACE1001311			47.002	40. 309	44.877	41.038	46.429	45. 578
20	PLACE 1001323	85.671	92.960			38.177	31, 146	36.745	54. 898
20	PLACE 1001325	63.854	83.048	40. 238	34.763				34. 275
	PLACE1001340	50.316	43, 105	32. 357	18. 188	41.779	27.080	44. 703	
	PLACE1001344	21.096	20.141	12. 901	11. 211	11.242	13.229	17.699	15. 374
	PLACE1001351	21.665	30. 334	17.172	16.561	21.087	13.674	23.521	25. 699
		51, 121	41.493	20.763	22.794	22.644	20. 945	39.950	30. 512
	PLACE1001366			8. 199	6.635	10.878	8. 266	14.816	8. 211
0.5	PLACE 1001377	17.643	7. 950		15. 238	10.327	18.369	19.779	20. 881
25	PLACE1001383	19.371	31.320	12. 152				21.749	13.042
	PLACE 1001384	12.523	28.763	17.012	8. 145	10.197	11.093		
	PLACE 1001 387	74.695	38.816	24.690	18. 993	17.630	44. 878	42.628	24. 984
	PLACE 1001395	16.685	20.986	21.294	11.232	11.885	13. 388	11.627	17. 398
		226. 500	168.857	120.411	105.668	74.590	106.559	109.855	113.693
	PLACE1001399		22. 276	6. 559	8.709	5.336	6.428	17. 374	13.590
	PLACE 1001401	7, 198				14.205	17, 551	44, 441	18. 269
30	PLACE 1001407	36.871	35. 435	20. 290	26.813			38.368	22. 732
	PLACE 1001412	37.695	27.537	14.076	15, 165	12.728	15.789		
	PLACE1001414	217.145	130.533	105. 385	81.994	74.052	115. 387	103. 177	72.729
	PLACE 1001416	35. 223	39. 103	34.029	25. 498	14.222	24.743	21.597	25. 005
	PLACE 100 1433	145.429	164. 813	104.366	153. 159	55.364	82. 221	118.995	143. 644
			39. 255	26. 807	18. 655	18.643	29.783	43.995	27.882
	PLACE 1001440	58. 228			47.264	46.872	43.771	53.047	50.036
35	PLACE 1001456	45.774	64.005	62.545		7.643	10.634	21.002	14. 923
	PLACE 1001464	14.904	12.569	12.016	7.606			17.688	12.134
	PLACE1001468	12.628	13.185	11.183	8.049	10.407	10.393		
	PLACE 1001484	111.986	88.704	61,951	103.045	57.131	47.838	72.549	64. 633
	PLACE1001500	112.534	66. 487	40.149	29, 195	31. 131	66.175	52.403	47.197
		111.530	51. 123	42. 187	21.773	27.041	52.421	55. 175	32.016
	PLACE1001502			47.845	42.970	37.990	50, 672	57.729	52.788
40	PLACE 1001503	104.144	79. 570		15. 526	12.841	22.000	19.770	21.944
	PLACE 1001505	20.479	27.535	13.492			18, 450	24.882	26. 311
	PLACE 1001513	30.859	24. 448	21.001	14, 991	10.141		71.012	63.994
	PLACE 1001516	133.217	89.711	99.042			73.719		
	PLACE 1001517	69, 164	42.110	28.024		19.564	24.584	39. 431	31.706
	PLACE 1001523	44. 322	26. 222			16.685	26.064	28.195	35. 152
	PLACE 1001526	12.214	48. 804				29.631	19. 107	33. 278
45			14. 916				22,739	18.831	18. 893
	PLACE 1001534	14. 278						19.891	10.823
	PLACE 1001536	25. 937	21.827				55.719	62.273	73.495
	PLACE 1001545	81.173	118.411						
	PLACE 1001551	29, 258	24.058	15,777	17. 582		14, 773		25.399
	PLACE 100 1564	12.683				6.981	12.704	27.781	11.258
				- 11			18, 406	16.691	18.288
50	PLACE 1001570	10. 554							
	PLACE 100 1571	127. 122							
	PLACE 1001595	116.778	213.788	32.313					
	PLACE 1001602	23.415		9, 921	11.848	9.736	11.310		13.830
		49.559					28. 595	39.306	37.052
	PLACE 100 1603								
	PLACE 1001608	26.740							
55	PLACE1001610	103.785	116.714	78.09	4 80.45	74. 242	31.490	1 05. 340	1 , , , , , , , ,

Table 129

		•		labi	e 129				
	(ALLOCHADICIL I	58.972	40.610	21. 168	17.897	20.458	26.980	37.282	29.415
	PLACE 100 1611		21.349		11.703	9.654	17.389	16. 943	17.712
	PLACE1001629		39.917	28. 058	32.960	30.608	36. 189	37.819	50.929
5	PLACE 1001632			12, 492	5.629	10.744	11.917	13.604	13.552
•	PLACE 1001634		22.871	23. 149	18. 271	16.901	31.188	43.749	17.679
	PLACE 100 1637	61.890	34. 286			29.961	30.898	32.648	42.726
	PLACE1001640	80.631	63.007	32.766	49. 291 14. 407	12.505	8.704	14.876	16.268
	PLACE 1001655	29. 386	40.949	10.818		19, 193	20.408	30. 495	28. 727
	PLACE 1001672	34. 615	40, 370	24. 145	16.896	5. 142	5.752	4. 884	4.020
	PLACE 100 675	10. 323	5. 349	4.889	7. 928		45. 917	57.578	71. 255
10	PLACE1001683		101.853	51.020	46. 928	31.257	27.851	24. 285	45. 922
	PLACE 1001691	55.061	48.826	32, 495	70.656	37.287	23.603	23.387	30. 475
	PLACE 1001692	50.688	45.778	29. 336	31.751	20.230	24.736	21. 290	26. 568
	PLACE 1001705	54.991	45. 920	32.949	30. 739	23.884		26.052	30. 073
	PLACE 1001716	19.961	39.584	17.983	14, 122	11.592	15.645		23.892
	PLACE 1001720	45. 804	36.576	23. 337	13. 159	14.367	26.395	38. 216	
15	PLACE 1001728	25. 294	12.023	10.018	4. 500	6.969	13.369	17.313	10.651
	PLACE 1001729	54.474	30.538	23. 378	14. 206	12.538	34.643	35. 119	24. 025
	PLACE 1001739	72.181	46.505	32.326	17.618	26.451	46.354	57.211	33.755
	PLACE 1001740	44. 321	37. 300,	20.706	23.395	18.627	20. 277	22.849	29. 188
	PLACE 1001745	88. 492	59. 243	42.077	24.655	33.811	52.589	78. 154	41.999
	PLACE 1001746	34.637	42. 251	39.371	25. 196	29.098	20.925	24. 039	30. 103
20	PLACE 1001748	68.976	42.569	32.885	20. 301	21.057	36.582	50.459	30.910
	PLACE 1001753	49.985	45. 870	23.560	22.075	3.690	25. 936	41.529	38. 920
	PLACE 1001756	58.884	78. 676	32.148	72.106	23.706	32.912	52.816	82. 360
	PLACE 1001761	80.396	70.047	114. 350	98.694	126.278	53.735	66.182	112.998
	PLACE 1001767	101.474	95, 179	45. 516	33, 144	52.766	54. 932	101.273	76. 611
	PLACE 1001771	19.712	26.759	20.057	12.622	18.385	15.780	19.880	23. 194
25	PLACE 1001775	4. 588	40. 521	8.311	6. 556	7.390	9.035	9. 583	17. 408
	PLACE 1001777	61.261	31.312	29, 820	13.022	17.840	32.541	34. 897	21.794
	PLACE 100 1781	16. 525	17. 889	7.311	9.028	3.652	9.892	13.994	12.461
		82.003	24. 962	30, 707	19.043	16.757	38. 137	43. 807	19. 485
	PLACE 100 1783	24. 406	20. 572	9. 992	12.368	9.648	12.063	27.946	22.791
	PLACE 1001786	39.981	29.419	23. 164	10.091	15.084	30. 627	38.055	36.556
30	PLACE 1001788	36.820	39.616	20.098	14.057	16.433	21.056	32.809	26. 943
30	PLACE 100 1795	128.712	38.515	26.836	13, 466	28.718	51,074	76. 434	36. 462
	PLACE 100 1799		17.039	10.361	10. 109	9, 092	9, 695	10.813	10. 585
	PLACE 100 1810	14.418	22.601	33. 584	11.211	34.814	38, 481	19.140	20. 248
	PLACE 100 1817	30.913	41.515	23.006	22.091	25. 640	19.095	24.750	27.083
	PLACE1001821	44. 377		20. 807	7.935	23.084	27,957	36.704	21.625
	PLACE 100 1836	51.521	27.558 29.744	21.870	21.220	18.464	14.961	23.954	18.459
35	PLACE1001844	29.459		18. 233	14, 133	19.354	20.298	32.062	33.894
	PLACE 100 1845	33.946	36. 421	15. 393	27. 399	23.094	20.179	32, 496	27.946
	PLACE1001858	36.762	28.558	16. 671	13.297	14.417	29.644	49. 283	21, 491
	PLACE 1001869	41.811	29.631	7.813	9.785	8.947	7.055	22.588	20. 287
	PLACE 1001890	21.015	19.216	18. 203	17. 788	18,625	34.484	37.521	38. 175
	PLACE 1001897	41. 587	86.444	26. 521	77.375	23.800	40.850	29.474	82. 496
40	PLACE 1001902	13.879	28. 323	18. 415	13.316	15. 185	24.027	48.664	25.843
	PLACE 100 1904	42.359		52. 221	54.031	60.482	55. 231	87.790	65.770
	PLACE 1001907	99.999	94. 157	33. 663	25. 331	33. 103	39.045	56.245	37.978
	PLACE 1001910	76, 138	126.370	43.604		51, 566	44. 297	53.061	61.896
	PLACE1001912	72.652	96.989			29. 328	46. 949	78.822	51. 365
	PLACE 1001918	59.029	60.982	33. 789		10,009	15, 594	8.844	29, 435
45	PLACE 1001920	9.437	24.354	8. 429				22.437	22.775
	PLACE 1001928	20.462	35.914	14, 995				22.090	19,274
	PLACE 100 1930	16.268	28.124	18.470					9.643
	PLACE 1001949		22. 587	13. 269			21.015		18.763
	PLACE 1001959		30.344	15. 913					16.621
	PLACE 1001969		20. 205	14. 372					21. 180
50	PLACE 1001974		45.767	37.839					21.729
30	PLACE 1001981		27. 300	20.961					
	PLACE 1001983		45. 469	30, 920					32.148
	PLACE 1001989		59, 400						60.144
	PLACE 1002004		138. 468						
	PLACE 1002 008			63. 83					
55	PLACE 1002015	48. 810	48.095	25.04	26.42	28.835	30.124	, , ,,,,,,,	
33		-							

Table 130

	PLACE 1002044	15.432	19.617	12.298	7.674	10, 740	14.882	16.986	23. 255
	PLACE1002046	35. 129	24. 586	16.894	16. 958	15.796	25.488	45. 998	25. 557
-	PLACE 1002052	13. 131	11. 184	10.040	6.082	7.542	10.153	10.668	10.355
5	PLACE 1002066	77.695	109.726	92.490	79.876	58.443	57.230	64.889	69. 207
	PLACE 1002072	97.971	90.711	48.605	48.732	39, 945	44. 244	40.362	47. 906
	PLACE 1002073	48. 101	39. 394	30.681	27.085	15. 219	30, 451	35. 202	22.863
	PLACE 1002080	147.011	90. 983	77.089	67, 438	53, 419	83.047	71.583	70.087
	PLACE 1002081	6.752	13.958	11.761	8. 303	6.211	11.142	11.382	8. 460
10	PLACE 1002090	19.854	27.734	20.058	14. 085	36, 381	18. 780	21.857	42.680
.•	PLACE 1002095	60.336	45. 829	29.642	33. 247	26.663	24.615	34. 539	41.411
	PLACE 1002 102	164.050	58.094	40. 254	32. 448	30. 279	73.576	158. 991 47. 534	75. 372
	PLACE 1002 109	45. 221	57.996	53. 572	43.855	38. 839 4. 013	7.023	5. 912	53. 651 6. 295
	PLACE 1002115	9. 512	11.954	8.778	7. 248 27. 115	43.709	26.254	23.542	33.029
	PLACE1002119 PLACE1002140	35.430 48.179	58. 455 44. 018	53.047 31.256	17. 883	20.743	30.803	35. 802	31, 498
15	PLACE 1002 150	14.549	14. 324	13, 952	8. 635	12. 089	7.434	7.940	13. 111
	PLACE 1002 153	99. 975	52.998	35, 156	18.899	19. 864	38.034	40. 428	32.754
	PLACE 1002 157	55. 938	35. 819	25.050	31.682	30.081	23. 109	34. 931	28.217
	PLACE 1002163	57.219	47.664,	19, 449	22.757	26.545	33.066	43.744	29.963
	PLACE 1002168	30.977	45.777	30.115	44. 322	21.088	30.717	33.746	25. 283
20	PLACE 1002 170	68.838	22.754	23.239	11.296	13.008	21.765	31.540	17.540
20	PLACE 1002171	23.819	23. 126	16. 254	25. 334	9. 191	13.358	14.604	12.880
	PLACE 1002180	18.621	18. 513	11.924	11.799	15.091	9. 384	14.450	16.442
	PLACE 1002184	11.237	16. 438	6.314	6.973	5.890	7.372	15. 552	5. 123
	PLACE 1002200	41.279	32.645	19.848	12. 160	14.612	26.495	24. 978	18.652
	PLACE 1002205	8.060	8. 833	8.840	5. 678	9. 502	7.453	5. 919	5. 027
25	PLACE 1002213	132.823	94. 631	54. 268 12. 888	62.752 18.583	37.757 11.494	66.436 15.981	72.589 15.553	61. 367 12. 757
	PLACE 1002219 PLACE 1002227	28.945 82.051	25. 808 55. 700	42.058	32, 436	34. 199	39.449	33.444	40.762
	PLACE 1002253	58.857	21.589	23.552	8.315	9, 457	21.335	22. 438	14. 348
	PLACE 1002256	11.668	27.097	12.608	15. 320	10.327	9.326	7.247	18.657
	PLACE1002259	12.944	16.713	14.115	16.119	13, 177	10.814	8.343	7. 436
20	PLACE1002285	12.935	14.107	10,661	5.670	8.397	8.906	13.661	8.898
30	PLACE1002301	40.882	61.873	38.880	19. 138	39. 970	34.344	28.064	32.685
	PLACE1002310	16.971	21.006	23.836	10.651	24.965	17.853	17.328	20. 350
	PLACE1002311 PLACE1002319	32.060 21.289	30.946	17.177	14.219	10. 905 9. 953	20, 580 15, 052	20.767 12.933	19. 139 13. 930
	PLACE 1002329	41.607	17.105 28.970	16.757	13.513	9.723	19. 282	28.768	18. 428
	PLACE 1002333	10.233	17.705	5.802	5. 259	5. 108	7. 829	11.050	8. 546
35	PLACE 1002342	48.414	46.073	26. 203	18.031	31.808	29.119	31.805	35.900
	PLACE1002343	38.774	31.024	21.839	9.918	13. 209	21.177	28.826	23.746
	PLACE1002355	37.547	27.979	16.049	8.792	11.795	19.972	18.057	19.576
	PLACE 1002358	48.964	52.954	25. 597	17. 560	25. 248	26.885	39.078	44.650
	PLACE 1002359	70.702	60.072	41.768	24.857	27.424	38.617	51.234	48. 247
	PLACE 1002374	119.415	70.407	40.003	52.366	27.254	71.202	86.975	59. 999
40	PLACE 1002376	76.607	80.189	66.224	38. 374	30. 440 8. 747	43.752 27.687	57.781 38.031	47.015 38.157
	PLACE 1002379 PLACE 1002386	45.960 34.135	37.677 56.039	24. 324	15.686 15.130	13. 263	40, 392	20.988	18.948
	PLACE 1002395	50.771	34. 342	21.705	12.792	17.447	30.904	41.999	26. 921
	PLACE 1002399	26. 369	26.554	11.941	11.546	12.821	16.487	21.773	21.163
	PLACE 1002407	24.383	13.800	14.460	6.932	17.857	10.390	8.160	9.349
45	PLACE 1002433	48. 909	60. 537	30.096	33.352	22.856	24.152	49.419	48, 535
	PLACE 1002437	41.702	30. 287	21.358	10.885	8.866	22.078	29.556	17.959
	PLACE 1002438	13.555	11. 187	8.617	6.781	2.684	9.005	7.945	7.896
	PLACE 1002446	21.605	27.628	11.792	11.569	10.494	11.830	17.464	16.893
	PLACE 1002447	35, 206	16.567	12.839	7.714	16.646	21.325	23.151 16.070	14.505
50	PLACE 1002450 PLACE 1002462	7. 279 28. 126	19.248	9.887	8.084	9. 639	12.889	28.071	18.658
50	PLACE 1002465	50.708	38. 829	28. 583	22.053	22.627	24.578	37.561	35.602
	PLACE 1002474	42.838	48. 831	28, 190	20.034	25. 208	37.936	39.355	29.560
	PLACE 1002477	68.476	88.049	43.373	49.594	28.828	30.662	33.024	45. 912
	PLACE 1002493	20. 932	15. 425	14.743	9.609	5.982	13.112	18.554	13.289
	PLACE 1002497	62.857	26.623	15.819	9.997	10.197	19.095	23.320	14.788
55	PLACE1002499	25.484	35. 975	17.658	12.207	20.785	19.603	26.553	24.711

Table 131

									45 400
	PLACE 1002 500	61.430	52. 592	20.851	20.792	20. 608	26.596	35.837	25.000
				25.761	14.063	20, 170	29.748	38, 465	28.873
	PLACE 1002514	57.950	34.821				15, 108	39.619	19.093
	PLACE 1002518	33.229	41.213	15.047	27.600	25. 421			
5	PLACE 1002529	20.589	17.020	8. 550	4. 795	6.064	5. 232	8. 483	8.689
•			81,-188	-71.765	41.993	49, 408	124. 500	121, 100	70.493
		228.966				32.704	37, 719	49.674	44.065
	PLACE 1002536	54.940	104. 532	50. 236	37.932				
	PLACE 1002537	50, 443	35.983	26.347	14. 124	16.394	28.846	22.586	18.551
		43.269	40.064	22.458	15.887	20.345	19, 917	47.789	34.032
	PLACE 1002539					32.445	28.657	42, 402	32.824
	PLACE 1002547	56.046	40.874	34.045	20.245				
10	PLACE 1002571	22.915	18.915	20.884	11.040	19.304	18. 369	20.827	18.977
	PLACE 1002578	110.554	134.909	53.782	65.675	56.576	47.716	58.650	75.950
					12.655	12.171	11.770	8. 242	11.466
	PLACE 1002583	10.726	15.813	12.765					
	PLACE 1002591	30.958	26.809	17.781	9.878	19.760	16.773	24. 345	16.337
	PLACE 1002598	14.446	15.092	4, 386	12.890	11.213	8.112	5.827	10.365
				19.490	23. 538	18.247	17.300	19.554	24.344
	PLACE 1002604	31.921	44.779						44. 841
15	PLACE 1002612	55.401	62.901	26.650	24. 921	30.069	38. 235	60.295	
	PLACE 1002625	23.240	23.910	6, 945	6.719	8.340	13.804	18. 338	12.847
					12.130	17.684	35.619	30, 109	30.357
	PLACE 1002638	47.938	43.765	20.041					60.920
	PLACE 1002655	99.112	95.019	46.543	45.871	43.662	48.343	74. 802	
	PLACE 1002665	56.436	48.910	34, 541	41.310	34. 121	40.016	45.653	42.518
		125.131	56.394	32.422	13.563	38.268	66.967	86.419	50. 297
	PLACE1002685				46. 294	48.882	52.021	80.560	61.182
20	PLACE 1002692	132.787	228.548	52.995					
	PLACE 1002714	44.319	53.609	23.573	28. 126	20.794	16.095	44.240	36.632
	PLACE 1002721	48.707	45.968	24.879	33.949	24.596	24, 407	47.991	34.094
			20. 165	11.297	10.959	22.220	21.294	29.351	14. 502
	PLACE 1002722	51.611				43.077	52.449	71.534	49.750
	PLACE 1002726	125.645	56.983	41.963	24.383				
	PLACE 1002756	76.684	90, 401	34.602	33.347	35.450	32.003	38.085	37. 112
05	PLACE 1002768	37.065	34. 695	22.471	18.473	10.495	27.644	30.569	9, 688
25					12.530	9.455	11.715	18,808	10.755
	PLACE 1002772	19.381	21.230	12. 133					
	PLACE 1002775	215.958	171.561	119,480	99.390	61.339	134. 546	191.663	118.381
	PLACE 1002780	176.781	287.195	23.632	43.077	19.593	82.890	72.700	18.752
			23.226	15.927	9.468	12.050	16.476	22.237	15.411
	PLACE 1002782	27.818				8.616	19.358	32, 122	23.951
	PLACE 1002794	34.691	31.569	16.222	15, 221				
30	PLACE 1002795	34.772	50.236	36.000	40.363	13.011	24, 050	29.340	37. 202
30	PLACE 1002811	40,778	28.219	23.615	10.194	9,406	18. 249	26.914	13.705
					9,929	13.556	19.575	20, 271	16.079
	PLACE 1002815	32.688	27.116	17.000				64.752	42.913
	PLACE 1002816	121.530	77.053	58.292	56.734	32. 151	78, 899		
	PLACE1002822	35.773	43.718	34.305	25, 631	11.831	23.639	48.755	30.733
		24.398	36.649	16.252	14.271	19.041	21,708	18.804	12.550
	PLACE 1002833					26.786	19.706	15.958	54.212
35	PLACE 1002834	20.377	29.028	18.884	38. 505				
	PLACE1002835	104.711	48.012	49.299	39.789	40. 131	89.778	70.476	54.471
	PLACE 1002839	22.755	19.054	13.353	10.924	8.604	13. 987	21.043	11.363
			22.474	16.954	12.287	11.607	17.683	15.934	14. 373
	PLACE 1002851	22.576				16.657	15.712	10.771	9.732
	PLACE 1002853	34.418	31.665	25. 145	13.903				
	PLACE 1002881	102.976	97.917	70.514	87.830	51.598	50.758	41.241	42. 291
	PLACE 1002901	71.548	63.698	66.555	29.645	45.140	59. 208	76.206	45.691
40			11.408	5. 948	6.331	4.476	4,773	15.458	10.017
	PLACE1002904	6.345					21.251	27.020	24.149
	PLACE 1002 905	43.777	43.201	24. 460		14.443			
	PLACE 1002 908	38.273	28.688	19.809	11.922	14.762	22.711	23.772	25. 263
	PLACE 1002911	280.363	142.219	110.57B		94.746	116.830	190. 264	121.060
					25.127	21.749	21. 182	28, 172	23.976
	PLACE 1002941	45. 141	51.204	25.368					27.392
45	PLACE 1002950	22.227	42.383	28.848			40. 551	30.415	
45	PLACE 1002955	118, 340	126.144	74.949	61.222	67.700	127. 593	138.479	103.622
							30. 246	30, 209	53.696
	PLACE 1002958	42.823			3.333	11.152			10, 215
	PLACE1002962	7.154							
	PLACE 1002967	62.925	77.879	33.266	40.751	36. 265	24. 991		78.774
	PLACE 1002 968	73.792					30.815	23.286	26.721
							_		
50	PLACE 1002976	24.111	38. 815	16.069					
JU	PLACE 1002991	83.434	88.462	43, 928	55.219	35. 522	33.200	32. 513	44.550
							27.082	27.450	28.611
	PLACE1002993	62.886							
	PLACE1002996	19.729	20, 547						
	PLACE 1003010	240, 363		98.21	60.019	42.226	129.379	119.840	90.413
	PLACE 1003025	68.787							
55	PLACE 1003027	22.588	27.019	12.98	10.960	16.947	17. 092	18.805	11.735
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Table 132

	PLACE 1003044	14.108	16, 171	12.882	10.168	11, 272	11.173	13.588	13.162
	PLACE 1003045	9. 931	13.537	6.830	5. 366	4.210	11.198	8.884	10.489
	PLACE 1003052	44. 591	46.375	21.677	18.989	17.471	26.652	30.514	25. 422
5	PLACE 1003083	20. 536	22.159	9. 236	10.342	7. 370	10.043	10.531	9.741
	PLACE 1003085	24.408	20. 399	11.964	14.547	6. 525	15. 327	21.584	12. 854
	PLACE 1003092	12.637	30.662	12.298	17.303	9.545	11.397	14. 192	24. 648
	PLACE 1003097	21.163	28.352	8.618	7.565	3.855	8.878	9.083	12.625
	PLACE 1003100	43.307	32.855	19.035	17.015	15. 982	50.024	32.500	18.851
	PLACE 1003108	58. 475	45, 704	33.791	31.380	26.209	26.815	25. 220	26. 126
10	PLACE 1003115	143.932	81,794	75.879	39.097	80.354	68.496	127.480	88.406
	PLACE 1003120	100.979	101.665	82.247	77.470	49.512	53.513	62.113	89. 513
	PLACE 1003135	6.556	10, 790	5.392	16.841	4,741	6.451	6. 382	9.459
	PLACE 1003136	55.512	44, 451	32.908	30.362	21.310	28.720	24.260	37. 347
	PLACE 1003141	7. 159	13, 191	10.628	9.244	4, 399	6.923	11.238	10.791
	PLACE 1003145	37.746	12.816	10.773	3.856	7.578	23. 487	24.678	15.744
15	PLACE 1003147	15.381	13, 149	11.750	9.884	10.068	7.642	10.640	10. 362
	PLACE1003153	70.554	49. 471	30.621	42.667	28.210	30. 997	31.700	41.448
	PLACE 1003163	37.733	16.360	12.470	5.123	13.824	40.304	50.483	17. 288
	PLACE 1003172	223. 164	104. 257	83.452	50.706	45.640	123. 594	116.341	107.613
	PLACE 1003174	6.847	14. 478	8.537	6.465	6.249	8.629	8.998	9.029
	PLACE 1003176	12.670	10,690	9.875	9. 192	3, 516	6.864	12.376	12.198
20	PLACE 1003181	11.687	8.674	6.252	6.507	4,411	6.989	5. 948	7. 466
	PLACE1003184	23.604	20.100	15.005	12.717	8.845	11.973	22. 555	14.655
	PLACE 1003190	12.444	5.722	6.366	11.024	5. 871	14.481	12.229	12.369
	PLACE 1003200	4.994	7.575	2.794	1.074	2.399	1.597	1.208	4. 980
	PLACE 1003205	156.027	157, 191	53. 553	83.830	63.878	61.050	52.411	61.365
	PLACE 1003209	19.507	25. 938	12.603	10.839	9, 269	15. 181	16.630	15, 534
25	PLACE 1003214	38. 350	83.164	20.591	69.513	15.776	19.528	39.872	125.749
	PLACE 1003229	49.722	43.024	29.429	25.068	15.677	21.087	17.077	23.421
	PLACE 1003238	17.754	10.174	7.246	3.501	3. 841	9.069	7.319	5. 314
	PLACE 1003249	51.840	53.347	30.500	32.695	22.004	24.099	28. 567	28. 591
	PLACE1003256	348. 304	244, 002	177.910	180.405	124. 873	188. 558	160.554	142.541
	PLACE1003258	11.993	6. 155	2.063	1.279	4. 364 63. 808	5.665	7.306	7.153
30	PLACE 1003279	141.943	126, 197	62.494	87.403		59.323 28.680	70.538 40.974	91.072 34.169
	PLACE 1003294	61.234	50. 989	24. 331	20. 131	23. 485 16. 935	42.888	30. 941	33. 241
	PLACE 1003296	41.072	45. 050	21.216	19.875	14.465	22.409	27.850	28. 987
	PLACE 1003297	21.895	44. 307	20.050	21.456 42.408	24.581	29.862	17.565	71.757
	PLACE 1003302	11.776	33.428 35.424	28.663 22.095	24.742	15. 104	19.475	23.808	27.587
35	PLACE1003334 PLACE1003337	28, 230 7, 957	26.706	3. 267	14.838	4.774	19.084	12.500	28. 263
33	PLACE 1003337	45.708	24. 591	13.442	10.821	11.910	22.698	29.220	24.007
	PLACE 1003343	17.266	13,753	6.616	6.894	8. 198	9, 061	13.065	7,734
	PLACE 1003344	323.950	233.808	153. 566	133.460	157.350	204, 264	266.356	264.565
	PLACE 1003353	53.698	66. 145	26.553	32.701	25.639	48. 208	44.219	57.187
	PLACE 1003361	84.141	102.796	46.744	55.344	40.194	47.082	41.263	49.755
40	PLACE 1003366	87.834	63, 858	27.852	28.427	27.117	31.747	33.446	27.075
,,,	PLACE 1003369	47.071	39.619	16. 521	17.558	18.957	16.856	24. 902	19.932
	PLACE 1003372	24. 973	37.849	16.679	21.014	16.249	20.971	27.530	18.337
	PLACE 1003373	94.491	102.178	34.895	57.049	44.893	39.537	40.009	45.753
	PLACE 1003375	36.319	27.954	14. 531	8.317	18.694	17.347	38.060	21.672
	PLACE1003378	10.936	9.134	3.801	3.628	4. 293	9.302	10.181	37.634
45	PLACE 1003383	23.472	30.580	11.017	13.956	16.293	19,925	21.999	14. 820
	PLACE 1003394	32, 582	51.968	30, 162	18.863	25.768	26.807	51.214	29, 166
	PLACE 1003401	24. 258	20:812			8. 433	7.409		10.841
	PLACE 1003405	200.792	69, 910	68.877		73. 544	91.798	149.248	62.838
	PLACE 1003407	150.376	60.878	43. 383		48.667	65. 167	94. 258	52.526
	PLACE 1003420	68.281	66.140	34.814		35.617	32.390	42.536	52. 238
50	PLACE1003428	34, 299	47.479	25. 133		23.830	14.848	52. 937	29.065
	PLACE 1003432	42.089	50.659	29.613		15, 118	31.218	32.711	33.577
	PLACE 100 3438	140.387	63.379	51.749		32.257	72.208	70.053	46.148
	PLACE 1003452	19.655	37.426	19.169		11.209	15.772	25.014	15.196
	PLACE 1003454	126.775	72.771	50.122		40.364	92.647	99. 924 96. 270	32.089 58.596
	PLACE 1003455	241. 296	81.923	63.513		58. 375 54. 057	56, 681	53. 136	61.523
55	PLACE 1003456	118.238	97.468	01.030	80.667	1 34.031	1 90,001	1 33, 130	1 01.023

Table 133

	PLACE 1003460	102.833	81.573	50. 363	49.760	31.621	74.863	91.750	61. 493
	PLACE 1003478	40.947	22. 624	17.515	11.339	9.308	17. 242	28.787	13. 341
	PLACE 1003484	93. 925	88. 475	99. 487	63.575	89.873	53.034	49.889	59. 266
5	PLACE 1003493		164. 272	115.044	85.931	75.866	116.655	174.628	103.683
	PLACE 1003503		147.014	62.133	98.370	49.594	72.492	61.425	119.480
	PLACE 1003505	22.557	36. 343	17.315	10.863	10.402	19.193	31.835	20.526
	PLACE 1003516	28. 486	21, 226	19.003	17.714	12.254	12.104	19.556	13. 167
	PLACE 1003519	139. 419	169, 111	84. 259	142.580	98.798	125. 181	74.459	170.077
	PLACE 1003513	122. 960	94. 921	139.217	91.721	132.495	50.021	63.727	115. 192
10	PLACE 1003521	17. 223	38. 437	23. 494	26.093	12.149	17.512	27.072	33.729
	PLACE 1003525	175.790	102. 294	111.400	74. 179	68.238	165.232	232.487	100.600
	PLACE 1003528	295. 594	306. 633		294, 409	135.971	193.013	253.930	588.036
	PLACE 1003529	198.617	81, 732	80.067	39. 906	48.188	118.411	106.875	72.890
	PLACE 1003537	25. 845	23.817	16.068	12.471	8.856	18.413	24, 516	16.865
	PLACE1003549	39, 079	30.714	21.959	18.378	20. 930	26.350	37. 243	23.667
15	PLACE1003553	44, 809	34, 386	16.950	14. 206	12.373	13. 162	26.918	17.820
	PLACE1003566	108. 286	89.542	60. 257	82.707	41, 128	52.437	55.865	<b>54.986</b>
	PLACE 1003568	19.139	24, 013	10.812	8.744	6.874	6.665	12. 296	6.869
	PLACE 1003573	28.529	30.963,	16.094	16.508	11.491	20.438	20.129	16.769
	PLACE 1003575	69.620	62.783	42. 283	48.323	24.844	28.963	18.823	28.675
	PLACE 1003583	13.478	10. 930	8.008	6. 298	3.054	5.089	10. 292	7. 945
20	PLACE 1003584	42.140	46.380	30. 421	29.764	19.273	18.780	16.951	29. 274
	PLACE 1003592	98. 964	131,059	76.620	85. 120	70. 369	47. 996	51.112	68. 235
	PLACE 1003593	2. 455	7.069	2.213	8. 879	4.615	4. 374	3.167 42.506	7.202
	PLACE 1003594	22.619	21.370	12.280	14.568	22.143	26. 231	17.988	29. 793
	PLACE 1003596	21.737	41.627	16. 247	20.950	11.333	20. 528 101. 281	103.552	70.110
25	PLACE 1003598	197.107	100.809	87.842	57. 151	53.833	15.616	14.765	13.631
23	PLACE 1003602	27.633	19.867	12.883	29.665	9. 628	29. 537	19.949	40, 233
	PLACE 1003605	13.132	30. 464	12.191 40.208	40.687	21.349	29. 193	33, 706	37.181
	PLACE 1003611	37.261	46.658	18. 870	19.849	15.708	16.877	16.986	14.810
	PLACE 1003518	22.786	32.124 16.778	13, 947	11.520	16.451	12.566	11, 487	11.026
	PLACE 1003625	16.924 94.235	146.631	108. 588	90.620	68.485	68. 227	76.568	114. 286
30	PLACE 1003626	66.350	38. 194	46.810	34. 410	25.790	40.498	47.259	42.273
**	PLACE 1003635	16.711	14. 910	11.010	9.614	7.598	11.425	11.967	16.224
	PLACE 1003638	42.833	38. 250	23.900	33.007	18. 152	21.521	20.597	26.688
	PLACE 1003644	32.340	47.319	50.856	35.086	25.956	21.602	39.503	41. 265
	PLACE 1003654	8.702	11.750	4.857	7.626	4.620	4.783	6.412	11.622
	PLACE 1003656	13.584	9.710	6. 305	3.192	2.939	12.859	10. 981	8.367
35	PLACE 1003660	48.712	57.359	34.700	32.321	18.997	26.256	31.653	37.509
	PLACE 1003669	18.575	20.941	11.934	7.933	11.712	11.416	9. 244	12.275
	PLACE 1003670	208.802	92,009	91.713	62.162	57, 129	95.305	111.081	85.224
•	PLACE 1003671	86. 484	44.662	34_415	21.211	22.436	39.601	73.903	47.170
	PLACE 1003697	20.072	30.957	22. 381	12.263	12.054	18.778	27. 550	30.714
	PLACE 1003704	37.863	72.473	31.653	35. 394	19. 396	26.513	28. 063 2. 305	51.598 18.174
40	PLACE 1003709	2.009	0.961	4. 994	3.081	0.994	36.509	43.017	28.963
	PLACE 1003711	69.991	36.386	26. 693 26. 163	20.921	23. 419	26.594	37.972	40.416
	PLACE 1003723	64.751	56.292 79.454	49. 180	55.077	46.271	43.499	54.124	55.073
	PLACE 1003724	108.825	29. 915	11. 983	5.933	6. 965	11.338	19. 286	17.493
	PLACE 100 3737 PLACE 100 3738	55. 859	28.082	23.047	12.820	11.647	24, 406	34.244	18.965
45	PLACE 1003742	45. 939	34. 288	20.111	23. 290	11.889	14.690	17. 236	19.752
45	PLACE 1003744	133. 197	117.135	50. 274	33.621	26.974	59.212	81.540	51.429
	PLACE 1003758	38.274	21.475	15.086	7.215		19.346	23.882	14.658
	PLACE 100 3760	26.760	76.015		18.973		50. 139	24.164	38.587
	PLACE 1003762	49.564	49.023				29.452	34, 554	31.123
	PLACE 1003765	85. 304	73.829				27.644	30.190	31.980
50	PLACE 1003768	44. 313	74.709				32.675		35.883
50	PLACE 1003771	21. 353	25.511						
	PLACE 1003772	15. 300					30.551	32.442	
	PLACE 1003783	21.327	19.915						
	PLACE 1003784	14. 398							
	PLACE 1003788	17.074							
55	PLACE 1003795	47. 580							
	L CV0E 1003133	77.300	73. 520						

Table 134

		. •	•	Table	e 134				
	PLACE 1003827	65. 231	45.890	25. 681	25. 602	26, 890	38.750	51.689	32.624
	PLACE1003833	108. 277			42.885	36, 400	47.382	71.347	50.389
		58. 333	54. 929		27. 226	31, 205	43.659	43.726	50.779
5	PLACE1003839	57.999	40.351		12.391	22, 439	35, 548	37.628	22.735
•	PLACE 1003845	127.357			23. 365	30.954	53.921	59.726	38.279
	PLACE 1003850	31. 154	57.883	11.655	9, 912	13. 233	18. 435	22.723	22. 101
	PLACE 1003852		20.593	18. 192	8. 183	16.756	18. 529	28. 232	22.273
	PLACE 1003858	41.915	44.111		13.711	17. 998	21.832	39.228	39.849
	PLACE 1003861	58.968	38. 407	12.279	15. 595	10.850	13.808	20.707	23.670
10	PLACE 100 3854	22.459	127.451	62.650	94.009	74. 206	51.275	57. 946	89.658
. •	PLACE1003870	60.423	33.558	22, 851	16.758	22.675	33.025	39. 475	20.419
	PLACE1003885	59.008	70.715	28.043	22.294	24.099	36.534	39. 216	36. 502
	PLACE 1003886	31.386	33, 156	12.296	8.686	9. 487	8.891	17.821	19.193
	PLACE1003888	9,030	10.854	5. 434	3.842	5. 628	6.081	11.548	5.474
	PLACE 1003892 PLACE 1003900	56. 299	34. 490	17.726	16. 257	24.111	27.255	40. 929	21.927
15		13. 429	29. 453	12. 159	9.597	16.000	7.119	13.508	10.317
	PLACE 1003902	42.879	27. 988	14. 980	9.315	15. 918	18.933	45.780	18.960
	PLACE 1003903	12.145	27. 163	12. 885	10.567	14.419	14, 179	19.072	14.872
	PLACE 1003915 PLACE 1003918	19.087	26.774	21.996	28.560	16.673	17.769	20.733	39. 272
	PLACE 1003918	17. 938	14.010	16. 114	10.428	12.304	11.045	13.677	13.626
	PLACE 1003932	12.148	25. 177	11, 239	11.640	9. 027	11.609	13.946	10.585
20	PLACE 1003938	98, 915	71. 254	54. 545	43.722	60.900	44.493	55.944	33.641
	PLACE 1003956	9. 602	25. 105	6.373	.20.612	8.000	8. 156	16.577	22.580
	PLACE 1003968	155. 632	59. 259	61.976	41.239	67.653	64.474	101.806	78. 393
	PLACE 1004018	54.312	58. 203	24. 249	21.023	23. 896	22.724	51.031	32.530
	PLACE 1004020	83.348	98.787	59.310	57.638	36.190	62.304	66.390	82.643
	PLACE 1004028	24, 781	24.415	11.783	9.512	7.540	20.059	22. 302	17.720
25	PLACE 1004034	17.910	20. 422	11.915	15. 479	8.400	11. 398	19.335	11.027
	PLACE 1004042	56.266	58. 516	27.953	28.385	23.656	36.706	42.138	31.854
	PLACE 1004078	55. 853	64.437	45. 957	50.360	28. 188	33.762	44.473	62.546
	PLACE1004103	82.183	108.065	67.258	73.844	58.609	52.188	48.497 42.824	30.092
	PLACE 1004104	28. 527	25.472	19.215	14.931	10. 266	14. 257 32. 925	49.556	52.027
	PLACE1004113	88.762	79.179	40.847	46.070	24. 179 13. 524	23.678	16.143	24. 982
30	PLACE 1004114	34. 482	51.070	24.001	25. 218	5.716	7.170	22. 166	12.479
	PLACE 1004118	7. 959	17. 781	10.600	6. 021 39. 964	32.057	70.811	86.636	81. 153
	PLACE 1004128	157.419	76.024	59.277	14.553	5. 947	9, 537	14. 197	13. 183
	PLACE 1004130	12.810	19.897	10.691	173. 146	139. 349	219.176	218.135	176.125
	PLACE 1004149	389. 247	289. 561 148. 253	89.024	112.406	77. 253	73.380	83.983	73.719
25	PLACE 1004156	154. 127 380. 298	97.742	180.381	64.718	155.863	307.172	350.794	93.857
35	PLACE 1004160	169.005	53. 952	58.840	40.858	55. 087	99.826	113.689	55.889
	PLACE 1004161 PLACE 1004166	34, 880	53. 232	25.983	22.633	19.476	13.270	20. 543	32.985
	PLACE 1004168	60. 294	31.301	30, 139	19.493	14.214	37.430	32.263	32.104
	PLACE 1004170	20.591	14. 931	15.171	7.631	5.880	13. 252	14.579	12. 294
	PLACE1004178	15. 161	17.955	94.893	8.136	138. 324	12. 187	13. 182	8, 196
40	PLACE 1004183	82.644	31.272	45, 235	14.851	34.570	64.117	65.703	24. 241
40	PLACE 1004197	16,554	15.430	12.063	9.295	4.620	10.632	9. 453	14.626
	PLACE 1004199	105.771	35.874	42.409	13.689	33.976	72.635	99. 795	29.738
	PLACE1004203	97.622	38.875	29.121	21.384	27. 473	49.790	59, 416	36.062
	PLACE 1004242	76.021	94. 358	69.979	49.223	46.887	46.304	63.046	50.392
	PLACE1004249		54.868	42.542	29.091	28.894	31.370	30.144	27.719
45	PLACE 1004255		6.797	5.218	3. 278		5.853	8. 367	4, 680
	PLACE 1004256	27.907	27.196	30.222			28.629	16.493	16.024
	PLACE 1004257		16.029	12,630			22.658	10.030	25. 329
	PLACE 1004258	25. 963	21.667	16.937			16.892	20.871	16.224
	PLACE 1004270	72.433	34.960	27.059			14.279	50. 286	21.577
	PLACE 1004272		17.600	17.337				91.578	15. 983
50	PLACE 1004273			38.516					9.115
	PLACE 1004274			7.313					29. 403
	PLACE 1004277								57.510
	PLACE 1004279								
	PLACE 1004282								
	PLACE 1004284								
55	PLACE100428	57.838	64.819	44.68	30.140	, , , , , , , , , , , , , , , , , , , ,	1		

Table 135

				lab	le 133				
	PLACE 1004299	72.960	55.550	24.647	14.840	20.355	39.409	45. 257	40.773
	PLACE 1004302	0.000	2. 283	0.000	2.351	1.896	0.000	0.000	2.398
	PLACE 1004305	48. 425	22, 731	21.012	11.875	13.056	23.176	27. 227	16.060
5		13.028	20, 460	9, 292	6.569	6.425	10.734	13, 111	14. 995
_	PLACE 1004316			3, 361	13.438	2,427	2.862	15.463	30.143
	PLACE 1004322	5. 597	14.420	97. 326	76. 987	66.867	109. 387	137.359	89, 195
	PLACE 1004325		142.328		95. 127	6,945	10. 855	19.856	120.861
	PLACE 1004332	20.898	80.056	11.714		74, 380	88. 121	64. 528	76. 583
	PLACE 1004336		109, 014	85. 745	88. 818		16. 208	18.604	15.045
10	PLACE 1004345	33.011	29. 261	14.760	16.784	13.075	151.939	184. 597	
10	PLACE 1004358	303.987	155.290	114.636	79.890	97.745		21.337	122. 929
	PLACE 1004376	26. 954	55.450	24.865	32.727	18.065	20.930		31.788
	PLACE 1004384	41.561	34.784	24.877	26.743	16.820	18.362	17. 481	21.709
	PLACE 1004385	2.815	8.008	1,116	0.789	0.276	1.941	4.609	1.615
	PLACE 1004388	9. 428	16.190	11.060	5.000	14.211	5. 122	9. 688	9.607
	PLACE 1004405	8. 173	12.654	5. 345	1.830	0.933	5.888	9. 305	5. 124
15	PLACE 1004407	29.905	23.442	14. 979	13.177	16.639	25.030	34.013	28. 941
	PLACE 1004424	10.514	15. 521	10.255	7.446	7.421	9. 255	10, 500	10. 362
	PLACE 1004425	19.759	20.897	10.508	10.323	4. 107	7.623	8. 354	13.395
	PLACE 1004427	27.135	16.966,	10.908	6.848	6, 240	12.886	13.355	13. 321
	PLACE 1004428	57.419	64.170	30.987	36.844	18.316	25.562	30.009	32. 786
	PLACE 1004433	14.267	16.470	9.620	5.755	5. 704	9. 109	18. 352	17. 362
20	PLACE 1004435	17.934	21.109	25. 397	11.056	16.381	15. 263	11.508	12.972
	PLACE 1004437	80.263	28. 301	25.518	11, 113	18.894	28. 285	48. 525	32. 402
	PLACE 1004441	54.134	47.973	28. 455	25. 980	23.238	32.602	42.800	31.312
•	PLACE 1004446	21.816	51.429	12.869	9. 278	14, 108	22. 134	27. 233	19.664
	PLACE 1004450	7.462	10, 131	7.421	5.906	3.377	6. 452	10. 209	6.066
	PLACE 1004451	20.207	31.572	19.505	19.989	13.665	10.206	15. 250	18. 302
25	PLACE 1004456	53.328	61.854	40.178	40.750	22.994	32. 386	43.215	51.423
	PLACE 1004458	11,625	26. 331	11.664	5.811	6.713	7.889	9.892	25. 194
	PLACE 1004460	14.565	10, 490	5. 224	4.840	4.848	10.082	10.381	9.064
	PLACE 1004467	55.048	46. 934	30.599	25. 322	18.898	22.765	24. 523	37. 228
	PLACE 1004471	79.809	63.442	37.258	59.178	37.277	29. 527	32.628	51.028
	PLACE 1004473	11, 959	24. 287	10.007	12.507	6.941	16.855	16.517	14. 312
30	PLACE1004475	28.089	59.714	31.110	18, 183	27.680	29. 310	26.516	47. 243
	PLACE1004482	25. 293	47.010	16.830	16.111	11, 400	30. 429	30.968	35. 155
	PLACE 1004491	1.664	6.234	6.645	3.270	2.102	2.892	5. 873	2. 357
	PLACE1004492	28.976	64.765	17.444	33. 197	14.425	13,718	15.087	46.827
	PLACE 1004506	115.632	78. 203	46.045	35.757	41.896	69.416	85.790	78.043
	PLACE 1004507	19.324	9. 642	5. 560	5.074	6.375	9.835	14. 279	13.049
35	PLACE 1004510	68.938	32.074	18.477	12.138	20.444	31.944	40.037	21.097
55	PLACE 1004516	12.480	28. 346	11.965	12.861	14.262	12.534	22. 485	21.487
	PLACE1004518	113.615	41.314	32.970	20.351	31.552	61.934	56.694	31.846
	PLACE 1004519	17. 977	18.444	5. 463	12.802	4.820	7.889	17.402	10.594
	PLACE 1004520	151.375	60.864	33.949	19, 465	34.865	66.695	80.040	24.602
	PLACE 1004530	43.149	50.004	13.982	11.859	13.432	25, 111	26.818	14.729
40	PLACE 1004545	10.167	15.345	7.071	4.082	3.066	9.778	48. 382	17. 084
40	PLACE 1004547	23.679	18, 172	11.002	9.917	9.918	8. 124	14. 641	11.578
	PLACE 1004548	65. 295	50, 486	25.299	24.808	18. 285	24.829	25. 884	36. 422
	PLACE 1004550	26.366	18, 052	12, 431	9.837	11.528	18.472	24. 539	12.011
	PLACE 1004551	36, 555	34.112	15.054	11.068	19, 459	22.324	30.835	27.019
	PLACE 1004559	7. 230	9,773	4. 555	3.840	5, 493	5. 484	6.749	3.314
	PLACE 1004562	28. 572	30. 296	23, 163	8.674	27. 528	15. 650	14. 237	9.875
45	PLACE 1004564	36.735	40. 092	17.343	20.204	19, 250	16.933	27.924	22. 272
	PLACE1004504	0.000	12.587	0.000	0.000	0.000	0.000	0.000	19.840
	PLACE 1004611	146. 180	120.698	55. 658	62.073	72.842	61.052	49, 103	61.906
					20.920	25.719	18. 242	25.782	34.340
	PLACE 1004629	33.357	43. 299	24. 243	11.469	40.312	43, 201	76, 589	27.684
	PLACE 1004630	115.833	50.627	40.441		25.704	57.715	75. 530	37.977
50	PLACE1004637	93. 560	57. 213	41.313			68, 433	63.089	99. 155
	PLACE1004645	73.214	93. 376	36.462		22.216	24,754	48. 692	
	PLACE 1004646	46.760	48. 123			15. 130			22. 337
	PLACE 1004648		101. 385	110.514			215. 200	161.060	64.085
	PLACE1004655	89. 992	149. 462	51.420		32. 385	132.613	125. 965	155. 546
	PLACE1004658		50.154						39.519
55	PLACE1004664		19.569	14.876	12. 928	9.845	14. 381	22.040	15.050
-									

Table 136

	Thi ACCIONALTO	115.072	106.617	82. 206	19, 303	40.425	71.021	72.226	74. 522
		31.963	33, 509	24. 678	21.646	15. 932	29.638	23. 353	28. 211
	PLACE 1004674		52. 263	26. 896	24.625	15. 862	23. 571	27.757	20. 193
<b>E</b>	PLACE 1004681	42.868	73. 361	53. 514	71. 286	30.833	39. 791	36.511	33.040
5	PLACE 1004686	77.947			14.534	30, 457	39. 275	32. 277	31.724
	PLACE 1004690	32.648	58. 935	35. 179		20.988	25. 836	29. 486	27. 807
	PLACE 1004691	54. 201	46.001	30. 198	29.746		10.119	16.032	13, 442
	PLACE 1004693	14.777	12.312	8. 393	5. 596	11. 162		35, 594	76. 510
	PLACE 1004701	70.824	100.375	71. 192	54.004	102.558	32. 216		
	PLACE 1004705	65.005	44, 191	23.752	22.321	16.770	23. 327	38. 083	39. 081
10	PLACE 1004708	27.110	53.686	25. 099	16.995	21.305	56.740	40.801	33. 150
	PLACE 1004716	39.167	36.771	27.872	31.814	17.418	21.095	22.468	30. 805
	PLACE 1004722	19.479	18.949	14.424	12.942	8. 398	12.638	19.361	14.771
	PLACE1004736	243.492	165.849	137.412	105.409	67.657	152. 337	156.408	125. 947
	PLACE 1004737	19.475	29.675	15.699	11.243	8. 873	11.625	22.792	35. 249
	PLACE 1004740	75. 304	51.308	47.454	36.445	39. 722	39.387	38. 438	45. 080_
15	PLACE 1004743	68.266	20.761	16, 980	15.277	16.469	24. 996	43. 820	20. 166
	PLACE 1004751	52.682	43.427	21.010	38.514	12.476	20. 526	37.750	28. 532
	PLACE 1004757	64.866	62.789	28.623	23.370	20.456	30.243	39.909	29.888
	PLACE 1004761	26. 949	16.825	13.926	8.696	8. 320	11.691	16.318	12.117
		54. 251	32.451	26. 443	19.663	14.012	23.566	35. 213	33. 476
	PLACE1004773	0.000	0.417	0.000	0.196	0.000	0.000	0.000	0.000
20		23.178	24.645	17. 477	11.418	17. 912	15.186	20.914	17.641
20	PLACE 1004777	10.099	9. 825	8, 108	2.235	6.900	9. 156	12.992	9. 524
	PLACE 1004793	188, 258	55. 088	53. 995	32.705	46.720	104.831	97. 648	39. 050
	PLACE 1004796	47.571	38. 570	28.854	17.511	18.650	30. 285	28.014	30. 229
	PLACE 1004804		19. 594	9. 102	9. 930	7.091	9. 407	7. 283	12, 102
	PLACE 1004813	13.617		65, 246	82.329	68.081	42.266	24. 121	54. 793
	PLACE 1004814	41.930	105.336		11.794	7. 165	7. 448	6.082	10.511
25	PLACE 1004815	11.260	11, 968	10.846	11.777	8.852	11.495	48.534	15, 257
	PLACE 1004816	16.128	75. 555		79.068	52.724	50.466	50. 930	68. 338
	PLACE 1004824	104. 392	119.714	59. 183	30, 150	21. 998	23.534	21.266	27. 294
	PLACE1004827	36. 438	26.140	22.831	12.887	15. 510	26. 557	30. 452	21.872
	PLACE1004836	31. 163	22.975	17. 358		19, 154	26. 439	33.316	30. 452
	PLACE1004838	51.513	33. 252	27. 542	18.538	4, 111	4. 374	5. 846	7. 493
30	PLACE 1004840	6.312	14.805	6. 440	3.431	12. 485	19. 399	19, 475	15. 636
	PLACE 1004842	36. 592	16.317	15. 880		14. 421	24. 250	37. 921	22.827
	PLACE 1004850	49.730	32.337	19.817	10.970	5. 213	6.832	14. 431	11.456
	PLACE 1004858	12.619	15, 190	6.828	7.862	13. 325	24.000	19.111	27.465
	PLACE 1004885	47.128	43. 214	27. 198		6. 285	8.607	7.712	8. 362
	PLACE 1004886	8.456	11.696	9. 985	10.337		29.704	27.795	64. 943
35	PLACE 1004887	25. 379	95.649	19.675	41.800	19.005	15. 543	32.352	19.012
	PLACE 1004896	15. 949	20, 476	11.823	11.627	11.685	67, 669	87. 798	52.760
	PLACE 1004900	156.735	97.505	60.889	55. 951	42.544		14.779	18. 931
	PLACE 1004902	34. 587	45.710	25. 541	18. 321	13. 921	16.696	11. 291	11. 270
	PLACE 1004904	13.083	9.418	10.864	6.532	3. 426	12.069		
	PLACE 1004911	9.050	2.555	6.611	0.560	18.979	5.276	77.886	87.866
40	PLACE 1004913	5,777	13.239	7. 908	7.304	5. 359	5.827	5. 467 6. 811	4.992 7.534
40	PLACE1004918	7.297	6.323	2.714	3.829	2.441	5.039		29.767
	PLACE 1004930	13.399	20.023	7. 288	16.589	5. 485	9.041	11.559 23.466	22.456
	PLACE 1004934	23.550	42.322	19.288	14.581	15.341	18.403	31.417	
	PLACE1004937	62.000	36.002	39. 437		29.690	26.536		16.660
	PLACE 1004949	54.760	253.300	30. 259	54.618	16. 463	68. 966	58. 166	114.761
	PLACE 1004969	34.833	23.924	16, 977	12.463	10.067	19.834	24. 891	18. 488
45	PLACE 1004970	0.656	0.020	0.000			0.298	0.381	0.000
	PLACE 1004972	6. 558	13.022	6. 101				11.774	11. 235
	PLACE 1004974	11.126	11.290				5.403	9. 800	10. 261
	PLACE 1004975	80.214	39.062					65.032	40.567
	PLACE 1004979	152. 165	104.604		83, 496	72.355	66.036	91. 372	96, 121
	PLACE 1004982	31.283	43.568	24. 303				20.250	25.778
50	PLACE 1004985	27.380	21.550	10. 343					9. 181
	PLACE 1005003	13.462	10.074	3.185					7.821
	PLACE 1005004	14.310	19.771	9. 570	8.29				11.577
	PLACE 1005005	68. 568		38. 586	41.076				
	PLACE 1005011	44.494		20.623	8. 452	15.065			34. 432
	PLACE 1005026	15.741					9, 113	16.290	8. 131
55									
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Table 137

			T	46 070 1	39.089	34.870	44. 104	36.457
PLACE 1005027		120.663		45. 870				36, 405
PLACE 1005031	53.784	80.972	22.926	20.892	23.652	30. 271	33. 677	
PLACE 1005036	59, 627	65,001	32.797	39. 527	17.608	26.473	31.634	38.145
LACETUUSUS		12.290	6. 164	5. 522	7, 108	4,000	7, 035	4.518
LACE 1005041	4. 201				38.790	39, 618	40. 595	41.015
LACE 1005046	87. 532	76.016	48. 856	61.696				23.815
PLACE 1005047	46.051	25.735		11.855	15. 156	16. 153	36. 409	
PLACE 1005052	46. 575	28.140	12.015	12.780	14.059	19.834	31. 197	29.860
		27.571		20.078	22.643	10.820	20, 439	26.659
PLACE 1005055	8. 158				25.138	25, 274	51, 837	39.544
PLACE 1005066	42. 175	53.415		15. 565				
PLACE 1005077	24.309	28.659	13.050	14.623	12.679	15. 734	21.504	21.488
	92. 222	93.468	34. 255	47.138	34.582	40.497	36.255	38.289
PLACE 1005085				57.228	50.800	42.000	46. 257	54.679
PLACE 1005086	102.289	115.876		31.220		196. 566	151.442	82.439
PLACE 1005088	544. 154	104.456		73. 371	168,988			
PLACE 1005089	15.670	20.631	11.122	11,637	9.823	8.077	15. 337	12.098
		118.635	90.799	64.835	74.093	133, 434	208, 569	89. 985
PLACE 1005101	240.793			67. 285	83.058	115.827	185. 343	115.880
PLACE 1005102	211.056	131.745	94.963				67, 557	
PLACE1005108	106, 691	120.848	45. 131	39.846	39, 785	42.063		31.335
PLACE 1005110	44. 564	38.347	24.937	14.829	19.447	30.115	34.784	22.848
			14.465	9.594	18, 283	14.066	20. 594	18.691
PLACE1005111	23.753	40.474,			43, 380	35.861	40.754	45.181
PLACE 1005123	59. 496	91.632	49.521	37.074				
PLACE1005124	40.401	51.742	18.340	18.486	14,709	15.661	58.570	27.105
	204. 940	150.075	112.018	89.631	91.526	103. 298	146.254	123.511
PLACE 1005128			31.043	64. 232	33.067	33.874	55, 788	78.228
PLACE 1005 130	60.815	73. 959			14, 502	14.628	19.090	38, 173
PLACE 1005141	31.384	66. 806	13, 194	14. 252				27.607
PLACE 1005146	41. 144	\$0.277	22. 100	13. 293	17.449	21.199	50, 528	
PLACE 1005152	24. 085	22,701	12. 226	17.968	9, 903	11.357	15. 172	18.599
		19.465	14.891	8.624	4.456	13, 395	11.532	13.083
PLACE 1005157	12. 965			20.024	12.111	12.771	17, 199	19,584
PLACE 1005 162	36.700	33, 286	16. 285	22.399				
PLACE 1005170	10.498	22.471	9. 375	11, 193	6. 555	8.512	31.001	12.095
PLACE 1005176	14. 622	9.067	7,477	7.780	4,490	12. 946	17. 364	10. 281
		9.688	13.589	5. 174	11,314	5, 046	10.911	5, 455
PLACE1005181	5. 793					17. 723	18.400	25.953
PLACE1005184	45. 108	51.852	28, 259	28. 577	14.895			
PLACE1005186	44. 227	18. 348	9,815	8. 521	7.622	25. 120	58.044	15.795
PLACE 1005187	35. 399	20.464	13.526	17.276	12.357	24.314	23.687	19, 988
		32.597	20,000	13.876	11.241	20.988	33.066	19, 839
PLACE1005189	22. 364			25.042	13.468	27.457	43, 397	28.759
PLACE 1005193	49.047	60.518	24. 364			40 007	36. 262	35.781
PLACE 1005200	33.619	67, 147	18.122	26. 564	10.723	25.057		
PLACE 1005206	7.546	16. 382	8,064	9.582	7.561	2.781	8.835	9. 588
PLACE TOOSE OF		12. 252	6. 329	7.983	11.377	8. 113	19.335	10.996
PLACE 1005216	12, 005			50. 792	22.094	32.500	31.112	40.20
PLACE 1005223	61.568	52.800	42, 403		13.973	30 303	34. 273	28.68
PLACE 1005225	56. 429	68.319	36. 547	41, 380				
PLACE 1005232	157.040	125. 455	69.019	54.944	48.079	58.072		47.854
PLACE 1005239	39. 974	13.868	24.220	12.450	8, 314	22.398	17.024	10. 214
			24. 574	15.713	15.164	30.409	32, 149	27.76
PLACE 1005243	44. 314	40. 194			9, 886	6.064		19.83
PLACE 1005250	16.580	27. 491	8.463	9, 418				11.28
PLACE 1005261	13.408	16.822	8. 222	5. 682	5. 972			
PLACE 1005266	20.535	27,721	31.380	28.026	16.734	16.639	19.888	14.31
		83.479	52,747	61.756	25.077	54. 250	44.786	57.87
PLACE1005271	93. 263			13. 425	7.075			12.24
PLACE 1005277	49.402	22.460	14, 621					32.91
PLACE1005287	22.199	38.345	37.586	27. 355				
PLACE 1005299	103.926	106. 254	44, 038	32.012	31.443		46.947	40.73
PLACE 1005305	31.910		25.573	14.702	9.928	36.933	23.937	7.78
	31.310			3.745				11.44
PLACE 1005307	8.172		16.098					18. 12
PLACE 1005308	40. 902	25.016	19.027	14.695				
PLACE 1005313	39. 342	24.175	12.571	9. 132	10. 374			21.75
PLACE 1005320	11,271			8.538		8.957	11.506	3. 50
								11.32
R. AZFAKAFAA-	17.688							30.18
PLACE 1005327		18.698						
PLACE 1005327 PLACE 1005331	53.315			23, 138	24, 128			30.3
PLACE 1005331			41,750					18.7
PLACE 1005331 PLACE 1005335	77.870	63.026				17,918	9.870	10.1
PLACE 1005331 PLACE 1005335 PLACE 1005336	77.870 21.324	63.026 20.435	19.530	20.249	15. 524			
PLACE 1005331 PLACE 1005335 PLACE 1005336 PLACE 1005351	77.870 21.324 322.456	63.026 20.435 95.522	19.530 98.703	20. 249 40. 129	15. 524 88. 620	198.28	224.069	67.7
PLACE 1005331 PLACE 1005335 PLACE 1005336	77.870 21.324	63.026 20.435 95.522	19.530 98.703	20. 249 40. 129 12. 918	15.524 88.620 26.29	1 12. 45	224. 069 8 22. 106	67.7. 17.1
PLACE 1005335 PLACE 1005336 PLACE 1005336 PLACE 1005361 PLACE 1005366	77.870 21.324 322.456 43.968	63.026 20.435 95.522 40.039	19.530 98.703 29.574	20. 249 40. 129 12. 918	15.524 88.620 26.29	1 198. 28 1 12. 45 1 30. 47	224.069 8 22.106 2 35.702	67.74 17.1 32.6
PLACE 1005335 PLACE 1005336 PLACE 1005336 PLACE 1005351	77.870 21.324 322.456 43.968 45.621	63.026 20.435 95.522 40.039 33.656	19.530 98.703 29.574 36.861	20. 249 40. 129 12. 918 29. 023	15.524 88.626 26.29 1 24.69	1 198. 28 1 12. 45 1 30. 47	224.069 8 22.106 2 35.702	67.7.1 17.1 32.6

Table 138

	PLACE 1005383	192.459	99, 179	41.513	26.019 [	36,659	74.701	68.796	45, 274
				17.165	2.620	2.702	5. 416	5. 640	1.066
	PLACE 1005388	13. 492	3.669	54, 915	55.853	33. 520	40.200	37. 456	42, 420
5	PLACE 1005409	90.786	74.023		13, 377	5. 674	22.632	23.974	18. 471
•	PLACE 1005410	46.290	42.715	17. 237		19. 395	33.771	55. 787	18. 201
	PLACE 1005426	91.681	34.075	23.596	8. 178	30. 455	29.753	21.758	27.288
	PLACE1005431	31,798	52.111	24.865	17. 489	41,772		21.743	44. 958
	PLACE 1005453	73.901	79.686	50.868	59.367		40.635		36. 491
	PLACE 1005467	53.538	58.699	26. 287	26.884	22. 037	19.003	24.688	
10	PLACE 1005471	14.111	22.568	10.718	9. 783	3.667	5. 561	7.986	9.086
70	PLACE 1005476	19. 213	15. 401	6,820	10.474	5, 214	8.066	10.246	12.895
	PLACE 1005477	44. 904	32. 541	21. 171	12.649	22. 905	16.973	12.374	11.640
	PLACE 1005480	15, 176	15.907	13.557	7.819	5. 374	9.674		
	PLACE 1005481	38.954	28.423	22.694	20. 287	10.897	21.409	20.874	20.662
	PLACE 1005494	3.769	10.339	4. 444	0.950	2. 290	3.620	3.635	4. 680
16	PLACE 1005495	66.511	51.739	18.659	10.826	24. 448	36.783	41.876	19. 394
15	PLACE 1005497	225.229	70.178	56.698	22.970	70.611	95. 227	102. 253	52. 394
	PLACE 1005499	34.460	64. 292	20, 603	24. 590	10.840	16.074	28.756	44. 984
	PLACE 1005502	23.366	16.975	25.072	11. 122	8.644	11.079	6.947	13.065
	PLACE 1005513	9.578	9, 101,	6.647	6.693	5. 372	7.954	6.929	7.661
	PLACE 1005515	26.055	17.913	14, 409	7.630	7.031	15.665	20.130	18. 654
	PLACE 1005519	1, 105	10.749	5, 162	20.785	2.814	7. 220	6.981	11. 525
20	PLACE 1005526	20. 332	17. 208	9.755	7.461	4. 693	10.134	18.343	11.671
	PLACE 1005528	135.917	114. 261	73.561	90. 213	64.605	59.074	53.101	76. 549
	PLACE 1005530	57.987	54.808	31.774	14. 143	29.079	35.603	50.048	45. 019 33. 023
	PLACE 1005536	45.147	63.002	37.450	8. 267	20.956	24.988	38.856	
	PLACE 1005539	124, 764	33. 255	11.994	7. 356	5. 220	14.637	17.879	10.020
	PLACE 1005543	44.082	34. 128	18. 253	25.879	12. 291	14. 141	13. 931	20, 699
25	PLACE 1005544	74. 900	40. 457	28.887	25. 245	13.758	39. 328	41.210	26.735
	PLACE 1005550	6.022	18.709	6.562	8. 947	5. 166	11. 247	11.859	13.763
	PLACE 1005554	12. 467	3.872	4.316	3.594	5, 956	4. 592	6.885	7.371
	PLACE 1005557	38. 341	19.894	13.342	7.004	10.123		24.623	20.113
	PLACE 1005563	49, 466	30, 178	12.647	9.014	15.593	21.940 30.056	32.864 27.968	27. 306
	PLACE 1005569	45. 144	91.673	20.105	17. 832	17.112 8.433	11.642	6. 292	7.748
30	PLACE 1005574	10.326	17.415	23. 239	15.035	1. 246	5. 392	10.776	8.407
	PLACE 1005584	1.575	8. 124	2.743	4. 127	8. 195	75.095	45. 627	11. 276
	PLACE 1005590	24.799	17. 304	10.072	5. 828	9. 204	8.707	25.759	17.524
	PLACE 1005595	23.048	17.414	15. 297	9.258	6. 390	6.373	13.351	11.411
	PLACE 1005601	19.725	11.146	9.146	3.038	7.570	5.089	9. 929	9.078
	PLACE 1005803	14.600	11.398	18.486	29.843	23, 139	24.076	25.335	30.827
35	PLACE 1005604	41.213	46.409	16. 274	16.607	8. 553	5.155	7.288	14. 586
	PLACE 1005611	8.443	24.450	10.537	8.137	6.368	11.349	12.772	6.731
	PLACE 1005622 PLACE 1005623	16.882	8.675 31.080	6.381	15. 139	12.715	20.665	16.500	16.140
•		14, 421 85, 952	39.001	28.845	20. 191	32.625	41.980	48.174	23. 375
	PLACE1005630 PLACE1005639	15. 544	15. 138	6.500	11. 153	7, 691	5. 800	12.445	10.861
	PLACE 1005646	77.577	49, 170	33.499	22.814	34.067	36.568	56.286	41.027
40	PLACE 1005647	24. 882	24.864	4.274	2.435	2.081	11.277	81.858	11.665
	PLACE 1005848	132.845	151.402	77.779	90.885	75. 286	60.577	62.598	76. 522
	PLACE 1005853	54.214	52.101	51.513	45.050	58.871	26.470	27.046	42. 423
	PLACE 1005656	10.886	10.384	4. 581	6.951	7, 146	4.012	9.841	4.680
	PLACE 1005659	66.511	28.923	22.280	14.717	20.121	25.706	37.588	18. 352
	PLACE 1005660	33.206	32.856	16.502	12.470	13. 584	17.875	18.205	12.323
45	PLACE1005664	111.456	61.079	40, 142	92.126	42.582	52.037	69.703	37.257
	PLACE 1005666	38. 297	57. 391	31.059		32.602	19.836	29.982	27.528
	PLACE 1005669	21.571	38. 576	14. 288		13, 912	15.528	26.157	24. 222
	PLACE 1005682	20.262	22. 251	10.868		10.729	18. 322	24.974	10.469
	PLACE 1005698	30.653	32.169	14, 400		8. 522	24.009	33.881	18. 345
	PLACE 1005708	70.622	71.219	28.705			39. 593	64. 431	43.104
50	PLACE 1005725	37.970	40.199	18. 153			15.434	20.139	15.072
	PLACE 1005727	10.738	20.546	10.308			13.636	6.798	18.026
	PLACE 1005730	31.961	20.066	19.504			18. 589	28.621	15.178
	PLACE 1005736	66. 424	61.842	32.233			36.600	35. 215	42.152
	PLACE 1005739	28.978	27.513	14.370		7.550		24.000	20.049
	PLACE 1005745	11.469	35.015	10.673	20.167	15, 864	28.058	24.092	16.459
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Table 139

	(AL LOF LOOK 753 ]	90. 237	41.210	18.989	8.672	12. 425	46.493	43.056	16. 151
	PLACE 1005752	1. 539	0.000	4, 104	1.918	1.510	0.000	5.784	2.632
	PLACE 1005755	66. 572	57.026		18.341	53. 529	53. 169	52.915	28.510
5	PLACE 1005756	79. 900	86.243	41.942	41.317	39.086	38.946	63.248	58. 527
-	PLACE 1005760	63.990	62.996	38.725	43.819	27.604	32.835	26, 439	25. 813
	PLACE1005763	118. 359	72.826	49. 483	36.802	35.749	50.090	71.856	51.056
	PLACE1005768	79. 421	64.882	40. 953	41.897	27. 292	23.749	34.685	36. 527
	PLACE1005771	37, 668	31.896	17.523	15. 262	12.345	17.985	18.238	19.301
	PLACE1005783	72.863	40.078	21.736	13.084	14. 828	29, 177	22.331	19. 278
10	PLACE1005799		17.722	27. 131	6.099	7.894	19.213	7.798	5. 528
	PLACE 1005802	6.212	61. 152	58.464	27.079	34. 644	91.079	90.094	47.378
	PLACE 1005803	16. 294	18.066	10.826	10.126	8.393	9.317	16.782	14.973
	PLACE1005804	75. 551	91.851	75.766	52.294	39. 477	54.790	85. 201	93.066
	PLACE 1005813	83. 027	75. 307	35. 260	46. 938	32.810	18.119	30.803	97.615
	PLACE 1005828	62.100	41.315	31.342	51.062	32. 258	19.627	15.080	24. 684
15	PLACE 1005833	15. 481	278. 446	15. 416	31. 374	13.721	24.043	14.331	47.385
	PLACE 1005834	3.601	10. 543	9.859	8. 251	9. 385	7.823	3.972	10.785
	PLACE 1005835	28.240	44. 997	17. 530	13. 182	10. 234	18.255	20.661	15.389
	PLACE 1005836	48. 952	28. 464,	13.401	6.803	8. 041	17.572	26.265	12.222
	PLACE1005845	6.922	14.049	6. 527	5. 977	6. 557	8. 274	10.956	10.665
	PLACE 1005850	60. 537	40, 486	33. 654	29.867	33.148	24.454	29.715	24.623
20	PLACE 1005851	5. 255	8. 502	7.076	7.967	6.349	5. 105	3.396	5.059
	PLACE 1005856	31.514	23.792	11.829	9.889	15. 184	17.753	16.532	9.402
	PLACE1005875	18.708	26.502	13. 111	7.247	11.323	7.852	8.071	10.929
	PLACE1005876	11.863	17.117	12.588	7.705	10.029	6.736	10.292	10.926
	PLACE 1005878	88.082	38.409	33. 471	15.538	10. 872	40. 432	40.415	25. 582
	PLACE 1005880	13, 768	23. 162	13.625	7. 279	4. 196	7. 444	9. 160	8.620
25	PLACE1005884	6. 339	23.822	4. 633	5.084	1.983	6.912	6.877	7.772 6.062
	PLACE 1005890	4.217	7.720	4. 562	7.386	4. 165	6. 206	4.379 30.359	26, 109
	PLACE1005898	49.218	42.891	38. 186	23.065	31.910	31.010 45.668	41.888	48. 362
	PLACE 1005913	88. 451	79.521	44. 625	46. 998	40.516	61.467	47, 211	132. 279
	PLACE 1005921	142.054	144. 941	38. 273	52.037	39.062 17.336	25. 033	14. 933	34. 055
	PLACE 1005923	63.053	60.900	27. 149	27.188 26.165	30. 860	26. 958	27. 906	18.684
30	PLACE 1005925	48.607	40.199 38.194	37.807 28.923	20. 495	16, 164	33.843	28. 337	44, 414
	PLACE 1005927	55. 705	16.013	5.744	4.478	1. 709	3.696	5.067	7.086
	PLACE 1005932	9. 087	56.236	26. 301	30.736	24. 397	28. 352	30.917	30.023
	PLACE 1005934	14. 496	14. 255	9, 508	3.415	8.672	4.033	8.619	9.076
	PLACE 1005939	123.849	544. 154	42.334	146.300	50.110	131, 268	94.038	701.375
	PLACE 1005951	30.248	32.418	15. 242	18. 690	12.128	15.271	23.652	24. 588
35	PLACE 1005953	19.693	12.970	10.718	9.877	7.414	11.462	12.609	10. 525
	PLACE 1005955	28.767	19.227	16. 323	8.434	5.041	17. 159	19.002	18.594
	PLACE1005966	12,530	5.651	4. 425	4.128	2.034	2.562	6.043	9.634
	PLACE 1005968	72.025	41.312	41.089	21.486	26. 270	41.994	52.960	35.566
	PLACE 1005975	25. 485	32.376	26.520	59. 431	24. 469	21.685	13. 392	59.446
	PLACE 1005990	28.041	21.763	14.040	6.899	9.815	14.633	20.007	16. 121
40	PLACE1005997	164.708	330.084	53. 780	239.364	63.798	139.506	181.530	287.794 62.761
	PLACE 1005002	107. 705	119.425	99,629	95. 897	48. 384	50.827	8.091	13. 582
	PLACE 1006003	17.046	17.747	14.438	8. 154	10.541	11.696 21.558	24.215	22.424
	PLACE 1006011	45.672	38.018	30.702	13. 512	11.110	18.839	15.505	19. 245
	PLACE1006017	45.647	36.734	21.158	25. 570 27. 384	13. 578	19. 303	24.570	28.170
45	PLACE 1006037	16.895	39.112	14.980	24.327	21. 931	28. 327	32.651	26.980
43	PLACE 1006040	46. 354	36.477	13.887 45.048		32. 191	49. 881	45. 260	38.790
	PLACE 1006063	93. 783	71.598	13.892		12.019	30.377	49.850	20. 945
	PLACE 1006071	21.534	36. 297	30. 172		29.043	23.961	31.954	27.041
	PLACE 1006073		57. 305	16.076		10. 251	15. 582	20.631	17.603
	PLACE 1006074		27.005	16.508			9. 947	9. 203	13.977
50	PLACE 1006076		32.791			21.503	45. 204	56.632	21.554
	PLACE 1006079		38. 429	26.815				13.690	14.363
	PLACE 1006093		15.385	13.757			15.676		10.524
	PLACE 1006 116		6.904	15, 512					12.644
	PLACE 1006119		12.984						
	PLACE 1006129								
55	PLACE 1006135	91.126	109.499	1 34.40	1 33, 033	70. 711	1 -200		

Table 140

	(Di ACCIONELA)	46.098	37.379	20.702	25. 574	15.236	19.435	15.116	22.985
	PLACE 1006143					5, 712	12.370	15. 306	9. 581
	PLACE 1006157	13.931	16.377	3.826	8. 200		12.185	22.648	13. 267
_	PLACE 1006 159	9.858	20. 502	51.646	6. 722	44. 269	7. 194	8, 960	10. 950
5	PLACE 1006164	16.798	16.274	7. 126	6. 999	8.372		95. 636	56. 466
	PLACE 1006167	167.052	67.298	52.083	32.075	42.820	71.882	30. 851	21. 127
	PLACE 1006170	53. 027	29.665	19. 393	10.419	17.774	25.072		14.042
	PLACE1006181	18. 281	16.157	7. 996	5. 350	25.260	14, 174	22.113	
	PLACE1006187	8.548	3.516	0.000	6. 582	4.888	7.985	6. 385	7.927
	PLACE1006195	29. 846	28, 480	17.352	16.886	13.459	17.228	26.763	6. 398
10	PLACE 1006 196	61.991	49.016	26.372	19.718	27.710	39.072	35, 118	29.050
	PLACE 1006197	54. 536	37.850	28.958	22.575	23. 293	25. 482	39. 927	23. 557
	PLACE 1006198	28. 596	28.607	16.575	17.769	13.452	15.976	27. 459 4. 973	22. 547
	PLACE 1006205	6. 745	7.609	4. 565	5. 214	6. 572	2.590		5. 687
	PLACE 1006208	27. 187	27.254	9, 873	14. 328	12.512	21.992	19.863	18.823
	PLACE1006211	51. 907	59.414	30. 208	13.725	28. 133	32.360	44. 159	27.440
15	PLACE 1006219	23. 493	24.408	16. 455	9. 362	17.274	26.290	25. 586	21.714
	PLACE 1006223	68.934	18.764	11, 909	9.616	10.504	6. 495	11.267	10.706
	PLACE 1006225	11.501	12.439	4.415	6. 582	5.792	8.314	14, 745	11.878
	PLACE 1005236	6.977	12.900,	5, 853	11.342	12.529	5.324	7.920	11. 191
	PLACE 1006239	22.381	23.765	14.765	10.878	15. 210	13.043	19.412	11.809
	PLACE 1006245	22.376	34. 520	10.634	11.051	12.665	11.374	19.724	21. 305
20	PLACE 1006246	7. 382	13.028	11, 301	7.187	12.507	7. 382	11.506	12.804
	PLACE1006248	26.428	39.894	16.473	21.809	14.977	13.745	20, 862	22. 348
	PLACE 1006262	31. 261	23.190	19. 574	15. 195	26.025	14.627	19, 352	15, 266
	PLACE 1006269	24. 853	29.569	14.626	9. 583	8.703	14, 129	23. 157	18.545
	PLACE1006275	102.949	70.174	48, 183	23. 852	33.229	45.824	59.434	33. 371
	PLACE 1006277	48. 240	62.171	21.255	16.104	9. 445	23, 300	38. 264	21. 261
25	PLACE1006288	70.893	32.184	31,657	17. 185	23.905	32.558	35, 514	20.818
	PLACE 1006290	10.445	14. 155	12. 302	10. 566	8. 624	8.747	18. 914	10.719
	PLACE1006298	31.578	46, 118	32.460	28. 976	15.993	23.096	26. 422	37.543
	PLACE 1006311	10.845	53.957	4.561	9.947	4.631	5. 498	6.778	11.014
	PLACE 1006318	58. 445	16.244	19, 191	15. 551	8.313	29.532	32. 903	13.674
	PLACE 1006325	22.893	33.926	3.989	1.894	3.728	40.444	14.737	21.889
30	PLACE 1006331	8. 939	11.370	13.783	13.776	7.560	9, 956	11. 998	18. 468
	PLACE 1006335	32. 529	28.387	14, 713	11.425	11.019	17.865	33.894	21. 152
	PLACE 1006357	3.825	9.950	6, 210	4.159	6.022	6.747	7.754	5. 087
	PLACE 1006360	14.089	16.595	24.796	8. 248	22.949	14. 298	13.022	11.859
	PLACE 1006364	50.974	44.777	21.918	23.821	14.219	27. 483	47. 224	26. 302
	PLACE 1006365	13.302	9,969	13.635	9.061	14. 422	9.214	21.696 34.207	7. 466 27. 153
35	PLACE 1006368	46.065	73.155	26.650	24. 050	13.240	24. 936	14. 801	7. 990
	PLACE 1006371	34.894	28.248	11.313	5. 383	9.407	18.791	19.815	15.474
	PLACE 1006373	37, 194	28.331	21.043	14.199	14.482	4.374	23. 912	14. 924
	PLACE 1006382	21.094	19.698	15. 454	9.638	8.482	36.061	46.518	25. 400
	PLACE 1006385	81.993	38. 251	25.850	13.853	17. 987	15.718	29. 802	21.518
	PLACE 1006391	24. 937	39.657	15. 251	12.115	12.857	40, 171	51.400	52. 505
40	PLACE 1006412	92.185	81.544	52.558	67. 133	2.933	4.693	8. 944	10.166
	PLACE 1006414	22.869	15.684	6.974	8. 725 15. 038	20.825	26.734	24.227	27. 471
	PLACE 1006419	61.800	27.143	19, 239	20. 259	23.756	34. 334	48. 209	27. 402
	PLACE 1006438	82.798	38, 554		67.312	72.074	128.015	104.908	86.500
	PLACE 1008 443	215. 537	110.762	106. 123	8. 147	6. 187	5.719	13. 324	13.219
	PLACE 1006445	11.757	18.560 37.610	10.002 21,247	25. 976	17,672	52.681	107.122	22. 397
45	PLACE 1006447	27.394		9.777	6.348	6.589	37.897	68. 487	10.963
	PLACE 1006466	16.826	15.029	<del></del>		24.572	43.598	56.094	27.697
	PLACE 1006469	114.915	41.384	25.605	34.721	19.002	28.695	34.080	43.083
	PLACE 1006470	55. 482	77.949		34. 982	34. 443	50, 263	43, 401	25.783
	PLACE 1006472	28.012	90. 945 54. 658			18.685	19.667	20.505	28. 511
	PLACE 1006475	82.952				21.907	16. 121	16.707	16.335
50	PLACE1006482 PLACE1006488	37.848	28.214 75.446			33.400	45. 132	55. 401	62.770
		97.835	112.335			45. 198	37.895	64.975	45.897
	PLACE 1006492 PLACE 1006506	97.220	13.735			10.467	11.563	6.929	9. 994
	PLACE 1006515	8.615				11.469	8.981	15. 280	14. 480
	PLACE1006516		17.795					13.079	19. 137
	PLACE 1006520	38.963	54. 680					19.311	31.254
55	FLEVOR 1008350	10.303	1 34. 000	1 32. 230				······	

Table 141

	In Act Abbed	76 630 1 1	03 128 1	42.948	44.567	33.031	39.882	33.174	40. 181
	PLACE1006521		03.128				10. 529	24. 356	58. 315
	PLACE1006529	53.118	57.618		32.693	19.830			22.594
	PLACE1006531	40.054	29.614	19.743	13.919	11.061	28.487	25. 077	
5	PLACE1006534	14.806	14, 541	8.631	12. 208	7.086	10.456	12.140	35. 132
•		111, 144		65.687	62.909	47.508	47.210	44.007	49. 020
	PLACE 1006540			33.934	19. 913	34.720	68.368	52.699	40.656
	PLACE 1006549	105.750	35.667				25. 245	30.050	25.681
	PLACE 1006550	53.734	37. 476	23.619	17.863	13.277			
	PLACE 1006552	36, 731	63.851	24. 515	30.033	16.150	29. 038	26.902	30.874
	PLACE 1006557	59, 138	32.373	20.742	27.767	14.998	53.010	66.775	24. 301
10			25. 131	12.554	16.291	12. 325	21.067	6.774	21.632
10	PLACE 1006563	12.150				11. 202	30. 323	25. 161	17.624
	PLACE 1006579	42.172	33. 427	19.515	12.744			10.056	19.854
	PLACE 1006594	21.308	62.751	8.959	11.953	18.053	24. 751		
	PLACE 1006598	38.010	39.953	22.806	22. 256	14. 136	17. 359	14.218	22.463
	PLACE 1006607	29, 363	43. 175	35.099	25.311	27.168	25.817	24. 362	33.010
		70.554	56. 140	32.568	26.861	32.156	41.824	78. 456	52.641
	PLACE 1006610		84, 729	48. 211	42.137	41.400	36, 165	33.872	88.891
15	PLACE 1006615	66.799				10.809	19. 146	13. 632	19.570
	PLACE1006617	46.945	34. 203	20.650	24.016				17. 583
	PLACE 1006618	12.467	22.675	10.936	4.988	6.177	12, 939	14. 170	
	PLACE 1006626	28. 824	22.724	12.096	14.424	6.491	15. <u>673</u>	20.994	16.846
		20.658	24.647	17.715	14.296	9.444	12.543	13.794	16.993_
	PLACE1006629			28.310	36.165	26.370	22.102	23.886	38.003
	PLACE1006637	66.078	44. 385			2. 901	12.736	2.835	3. 364
20	PLACE 1006640	1.906	3. 182	1. 497	1.850			40.050	24.555
	PLACE 1006644	47.828	33. 193	17. 215	13.059	19. 569	23.838		
	PLACE 1006657	19.786	8, 124	12.247	4.403	6.268	5. 198	7.763	6.121
	PLACE 1006673	45. 242	43, 900	31.743	33.164	17.416	21.697	21. 275	29. 264
		16, 105	18.660	7. 229	6.676	2, 905	9, 955	9.738	6.953
	PLACE 1006678			64.876	54, 439	35.908	50, 796	60.748	73.192
	PLACE1008682	108. 821	86.487			2.823	4, 669	8.079	6.963
25	PLACE1006584	12. 327	5. 526	1.745	4. 542			21.757	18. 483
	PLACE 1006698	35.079	26. 331	16.481	11.898	16. 188	18.313		
	PLACE 1006704	86, 472	27. 708	22.553	11, 168	23.040	31.772	42. 206	22.041
	PLACE1006708	63, 065	64.979	29.269	36, 158.	32.310	29,740	35. 534	34.620
		83.669	46.735	36.469	20.073	24.293	44, 745	40. 284	31.562
	PLACE1006711			19.709	9.911	12.634	19.601	15.694	11.421
	PLACE1006714	24. 897	21. 232			9. 686	25.065	16.540	13.432
30	PLACE 1006716	43.488	17.230	9. 950	6.619				19.367
	PLACE1006731	28.782	29.180	22.410	16.665	26.985	19.586	12.657	
	PLACE1006754	36. 921	20.331	16. 512	14.887	10.304	20.093	26.461	37.338
	PLACE 1006760	37, 757	42, 174	22. 283	15.705	21,554	21.150	17.013	41.393
	PLACE 1006779	3, 647	8.616	3. D16	6.280	6. 191	5. 298	7. 122	6.796
				38. 409	19.483	30.410	47.327	64. 324	35.890
	PLACE 1006782	92.507	28.870			10.900	18. 396	16. 357	16.765
35	PLACE1006783	27.658	31.732	12.496	14.567			15. 312	7.438
33	PLACE1006786	24. 498	14.495	7.472	4.210	11.343	13. 380		
	PLACE1006792	77.449	84. 545	47.367	55.539	38.143	39, 428	24.476	35.695
	PLACE 1006795	9, 133	4.460	1.737	2.793	3.353	3, 139	2.968	3.320
	PLACE 1006 800	4.005	5.373	6. 293	5.585	5. 488	3.372	4, 355	6.632
			18.118	5. 886	6.406	8.461	8.216	2.942	9.555
	PLACE 1006805	10. 412		18. 294	14.933	24.393	18.264	52, 680	31.248
40	PLACE 1006809	42.846	42.011			18. 598	11.836	22.056	24.307
40	PLACE 1006815	28, 382	27. 387	16.127	14,696			4. 844	0.000
	PLACE1006819	2. 234	8.095	0.000	2.742	7.006	3. 430		
	PLACE1006820	88.654	108.172	51.115	52.888	36.795	40.511	48. 278	48.233
	PLACE1006826	36.400	44. 215	19.975	9, 428	19.371	14.819	20.833	17.598
	PLACE1006829	92.548	43.863	26.240	21.591	27.592	41.457	58.358	33.442
					51.037	13.795	25.742	31.212	23.318
	PLACE1006853	36.698	17.968	19. 226		4, 197	4.805	5.604	5.225
45	PLACE 1006860	6.034	4. 924	7. 203				22. 604	24. 184
	PLACE1006867	38.603	40.857	22. 938		24. 586	16.186		
	PLACE 1006875		34.942	8.578	8.800	8.892	8.348	13.170	11.720
	PLACE 1006878		23.697	15.013	10.894	12.955	15.847	22. 292	15.804
				32.894			25.744	33.055	31.151
	PLACE 1006883		68.499						10.343
	PLACE1006898		7.894	4. 988					
50	PLACE 1006 901	21.369	32.566	11.362	7.983				15. 803
	PLACE 1008 904		60.723	40. 359		22.863			30.368
	PLACE1006917		18.119	4. 506				14.762	18.898
	PLACE1006932		50.295	37.537					
	PLACE 1006935	26. 622	22.255	28.03					
	PLACE 1006 956			13.80	17. 258	7.757	23.631	21.753	16, 324
55									

Table 142

								11.106	
PLACE 1006958	24. 224	20. 988	2.886	4.740	6.547	12.414	18.682	11.106	
PLACE 1006959	18, 928	26. 190	17.859	8.749	10.471	20.650	31. 538	10.229	
				45. 983	28.766	40. 349	60. 294	45. 326	
PLACE 1006961	117.650	81. 345						25. 186	
PLACE 1006962	45. 285	44. 217	26.483	25.012	20. 091	22.762	20.963		
PLACE 1006966	28. 233	14. 490	13,064	8.732	12. 926	14, 261	20.842	9. 575	
		17. 092	9.075	8. 221	7, 276	14. 248	14.630	10.668	
PLACE 1006979	17.727				7, 697	14, 839	32.067	28, 756	
PLACE 1006989	32.865	52. 943	17.860	11.639					
PLACE 1007001	63, 189	31,010	16.872	11.652	13. 459	33. 428	28.562	26. 941	
		49.098	38.389	21.381	19.097	50.704	38. 424	25.032	
PLACE 1007014	92.804				10. 271	12.863	18. 290	11.436	
PLACE 1007021	32.615	23. 234		10.544					
PLACE 1007026	6.113	17.016	5. 244	5. 923	5.797	4. 186	5. 493	10. 123	
PLACE1007028	32.763	23.055	16, 841	11.266	15. 159	13.728	15.276	14. 576	
PLACETUUTUZO				81.140	73. 181	232.551	242.218	764. 485	
PLACE 1007038		1311.392				14. 427	15.726	19.822	
PLACE 1007040	29, 591	22. 423		13. 542	14. 126				
PLACE 1007045	78. 257	39.847	30.671	22.858	23.390	22.928	15.061	21.566	
		2230.938		376.525	527.636	419.669	96.387	239.735	
PLACE 1007048					8, 179	14.765	16.384	14, 731	
PLACE 1007053	25.010	19, 115	11.205	9.097					
PLACE 1007068	99, 855	72.463	39.350	24. 132	16.753	40. 977	65.159	29.062	
PLACE 1007070	18, 155	27. 141	16.021	17. 985	10, 589	22.789	20.149	22.755	
				28. 282	20.816	24, 263	20. 952	30.465	
PLACE 1007076	36, 900	36.555	20.522				45. 495	14, 900	
PLACE 1007077	45.865	32. 193	19.090	3. 110	15.647	30.538			
PLACE1007081	5, 244	5, 196	3. 378	1. 304	2.199	3.337	3.589	2. 171	
PLACE 1007082	55.736	39. 537	14.678	18.774	16.347	23.656	45.049	21.718	
			7.344	11.776	17.009	14.076	7.700	7.525	
PLACE1007092	16. 389	10.500				24.039	25. 213	11.883	
PLACE 1007096	46. 332	24.876	22.197	12.502	9. 398				
PLACE 1007097	34, 116	23.336	12.085	13.012	5. 587	12.093	31.892	15. 157	
	57.957	45. 253	26.945	15, 165	13.161	35. 273	26.948	25.079	
PLACE 1007099				9. 937	8. 933	12.714	17.885	14,722	
PEACE 1007105	28.626	17.036	14.234				130.751	40.877	•
PLACE 1007108	41,006	85. 910	11.197	12.028	13.853	86.217			
PLACE 1007111	8.964	10.681	5.940	7. 255	7. 501	9.749	5.640	8. 886	
	30, 195	16.582	14.410	10.804	11.077	14.707	17,795	20.354	
PLACE 1007112					2.918	3.208	6.435	5. 903	
PLACE 1007130	11.359	6.838	5.607	4.816					
PLACE 1007132	68. 292	55. 387	61.678	43. 595	44. 456	42.578	73.359	40.514	
PLACE1007140	24, 801	47. 103	18.726	21.699	14, 109	24.706	33.892	29.052	
		21.700	13, 298	16. 396	7. 325	14,674	16.496	15, 455	
PLACE1007143	27.771		1		10, 493	14.373	24.878	12.622	
PLACE1007169	21.059	24. 932	10.043	15. 314			12.459	9.702	
PLACE 1007178	29, 316	18.952	15. 204	8.851	14.010	19.633			
PLACE 1007190	28.853	21.235	6.481	10. 255	7.822	10.991	13.037	15. 192	
	20, 919	11.754	12,200	7.867	9, 329	15,651	10.737	9.150	
PLACE 1007201				35. 864	26.097	42.107		71.342	
PLACE 1007202	75.891	83.211	41.376				20. 020	22.254	
PLACE 1007226	38.727	32. 391	24.013	15.641	12.748	28.566			
PLACE1007238	37. 920	27.260	52.707	11, 101	5. 882	19.768	19.683	17.554	
	25.792	17.879	12.822	11.697	11.572	18.220	21.634	16.456	
PLACE 1007239					7. 387	15.238	18,734	11.918	
PLACE 1007242	30.312	21.645	13. 524	8. 187			10, 310	9.966	
PLACE 1007243	16.786	6.525	8. 256	6.326	5, 657	7,341			
PLACE 1007247	47.743	24.409	31.744	16.238	32.693	32.792		21.768	
PLACE 1007257	50, 989		26. 453	23.676	21, 435	26.525	35. 446	30.498	
				45. 986	28.012	27,790		40.126	
PLACE 1007274	63.868	57.917	46.739	43. 300	10.012	21.277		25,771	
PLACE 1007276	45. 004	47.623	29.716	29.699					į
PLACE 1007282	51.770	26.821	22.456	16.571	9. 849			14.968	į
PLACE 1007286	51.312			41.722		28.174	21.307	30. 962	ı
				7.019				16, 151	l
PLACE 1007296	8, 691							4. 970	ĺ
PLACE 1007301	14.846								l
PLACE 1007314	170.251		56.463	38. 977	43.654	76.971	91.606	68.061	į
	7, 805					9,797		7.716	ļ
PLACE 1007317									ļ
PLACE1007329	22.649								ļ
PLACE 1007338	32.760	36. 157	17. 328	12.019					į
PLACE 1007342	35. 584					19.638	24.471	16.054	ı
								14.508	ı
PLACE 1007345	27. 643								ļ
PLACE 1007346	84.876	67.312	49.862	48, 124					ļ
PLACE 1007359				10.401	10. 905	16.783	28. 957		J
							58, 114	52.219	1
PLACE 1007367									1
PLACE 1007375									4
		32.95	18.430	10.509	5   15.018	23.30	0 20.623	14.853	
PLACE 1007377	44. 02	, , , , , , , , , , , , , , , , , , , ,	, <u>, , , , ,</u> , ,						

Table 143

				1000	10 201	6, 191	27.672	67.719	218.918
	PLACE 1007386	18.828	87.737		10. 203		4. 825	14.652	34, 452
	PLACE 1007392	8. 222	11. 434	10. 749	9.637	5.668			22, 559
	PLACE 1007402	65.708	33.760		10.518	17. 357	31.450	39.891	8. 437
5	PLACE 1007409	9.770	9. 329	3.971	4. 482	6. 413	5. 266	14. 242	
	PLACE 1007416	27.788	14. 552	13.712	11.561	17. 284	15.858	12. 261	17. 200
	PLACE 1007420	46.820	65. 531		15.727	22. 458	25. 870	29. 321	26.656
	PLACE 1007431	19. 972	36. 820	4. 499	11.250	12.525	8. 981	18.986	18. 539
	PLACE 1007450	45.777	50.126	22.855	30.226	21.905	18.828	17.972	24. 571
			46. 157	8. 675	25. 984	25. 596	10. 982	23. 901	19. 624
10	PLACE 1007452	33.958	122.886	31.320	44, 109	41.875	41. 307	59.818	59.212
10	PLACE 1007454			25. 529	18. 180	20.772	23,068	34.418	21.265
	PLACE 1007460	46.871	45. 449	12.040	19.617	18.742	17, 249	22.181	21, 235
	PLACE 1007478	30.938	25. 400		12, 842	15. 645	21.889	39. 282	17, 141
	PLACE 1007484	35. 483	18. 194	16.643	2.621	6.264	5. 521	13.139	9. 035
	PLACE 1007488	12.070	11.216	5. 905		10.052	11,006	19, 984	19, 609
	PLACE 1007507	16.065	19.266	11.755	10.003	5. 991	6. 407	13.848	8, 173
15	PLACE 1007511	12.031	9. 468	5.676	5. 965			25. 383	15. 457
	PLACE 1007513	28. 839	33.816	17.234	10.351	5.817	26.217		
	PLACE 1007524	31.989	52.731	17.490	18. 194	13.641	11.134	17.227	20.016
	PLACE 1007525	53, 144	47. 497.	20.989	29.065	21, 557	14. 406	17.969	19. 213
	PLACE 1007537	114, 162	62.590	29.450	28.798	42.322	39.868	74.479	42. 203
	PLACE 1007544	13.698	23.058	10.584	10.736	6.412	6. 388	19.809	12.059
00	PLACE 1007547	34. 533	43.022	15.777	19.820	14.818	10.460	22.065	27.725
20	PLACE 1007557	68. 240	54.730	20.858	22.219	17.520	22. 378	30,659	28. 149
			37, 749	42.477	13.441	33.714	29. 243	17.177	19.085
	PLACE 1007560	56.701 19.954	13.569	9.536	4.633	3.515	11.734	10.232	6.747
	PLACE 1007565			3.081	3. 286	2,111	3.703	7.852	3,004
	PLACE 1007580	5.661	16.015	19.036	4, 553	5. 377	21. 293	19.488	5, 045
	PLACE1007583	21.325	12.320	12. 980	17, 204	13, 786	15. 975	15,980	13, 540
25	PEACE 1007591	23. 357	23. 264			7.678	9. 978	10.043	33, 199
	PLACE 1007 598	10.914	22.683	12.140	19.473	0.000	4, 051	14, 144	7.161
	PLACE 1007610	8.777	5. 574	4.440	3, 931		9, 540	14.682	10.695
	PLACE1007618	27.729	17.405	12.198	7.493	7.679		21,828	41.949
	PLACE 1007621	127. 255	33, 162	30.450	23.070	22.170	28. 865		54.570
	PLACE 1007626	52.820	41.475	28.151	29.773	20.867	60.602	51.332	
	PLACE 1007632	59.751	36.549	27.076	16.433	16.357	35. 583	30.758	23.467
. 30	PLACE 1007635	54. 365	34.862	13.465	8. 465	10.812	17.884	31.723	24. 974
	PLACE 1007645	35.884	32.380	12.803	11.465	4.647	16.976	17.366	15, 901
	PLACE 1007649	22.119	4.188	5.061	14.689	4.509	20.917	21,502	5. 164
	PLACE 1007659	68.472	46.570	26.862	59. 476	24.769	18.505	25. 281	32.267
	PLACE 1007669	68.844	76. 485	26.431	38.944	24.278	27.709	17.065	31.698
	PLACE 1007677	35. 578	30.684	12.552	23.334	10.440	22.611	14.842	25.043
35	PLACE 1007688	56. 110	18.042	22.153	6.473	14. 256	12.150	17. 233	6.418
		6.860	17.051	10.688	8.318	11,590	6.899	7.099	22.589
	PLACE 1007690		6.551	4. 310	0.941	2. 439	6.854	5, 985	3,880
	PLACE 1007697	12.184 60.683	12, 143	7.740	2.796	6.156	6.869	11.415	8.331
	PLACE 1007702			8.512	4.274	16, 193	10.241	23.445	15, 595
	PLACE 1007705	40.045	12.817	11, 130	5. 527	15.086	10, 115	26.633	16.152
	PLACE 1007706	39. 169	33. 551		7.814	15.584	9. 357	10.094	10.940
40	PLACE 1007725	21.127	27.357	11.385	3.729	5. 453	10.931	14.086	2.233
	PLACE 1007729	28.499	11.383	5. 377		6, 592	10.898	20.320	10.820
	PLACE 1007730	24.859	34, 871	14.038	4. 450	21.728	24.240	17.956	20.227
	PLACE 1007737	64.586	44.554	26.554	35. 091		0.000	2.807	3.029
	PLACE 1007743	0.859	3.414	1.135		1.756		23, 929	
	PLACE 1007746	32.087	24.843	12.795		15. 204	23, 195	19.779	
45	PLACE 1007753	45. 192	21,910	9, 160					
	PLACE 1007769	10.061	8.971	6.218	3.760	4.071	5. 692	14. 415 29. 565	
	PLACE 1007780	67.441	127.130	21.733	15. 299		23. 156	29. 505	
	PLACE 1007791	23.878		11.597					
	PLACE 1007807			5. 484	6.978				
	PLACE 1007810	4. 996							
	PLACE 1007814						42.572	9.703	
50		27. 262						16.434	6.803
	PLACE 1007828								
	PLACE1007829								
	PLACE 1007841								
	PLACE 1007842								
	PLACE 1007843	5.632	5. 828	4, 884	2.27	5.807	2. 324	1.58	3 3.313

Table 144

PLACE1007845	3, 434	6.356	3.584	2.435	2.658	6.674	5. 597	3.658
PLACE 1007846	40.170	23.220	10.470	9.642	6.328	8.370	6, 111	14.996
PLACE 1007848	12.413	17.578		13.557	4.620	7.164	5. 426	9.910
PLACE1007852	3.936	5. 252	4. 986	2.146	2.510	0.958	1,562	2.641
PLACE 1007858	4.377	15.699-	-8.840	10.046	13.186	7.564	4. 439	10.001
	58.984	20. 206		15.579	24. 307	27.536	24.518	13.052
PLACE 1007866	204. 996	132.437	121. 332	60. 458	61.024	130, 512	134, 180	110.829
PLACE 1007871	75.858	20.469	18.620	9. 121	11.830	21, 548	25.804	17.607
PLACE 1007877	15. 982	20. 582	3.622	6.710	3.655	14.406	12,913	15.821
PLACE1007878		6.128	4. 453	3.005	1, 236	3.305	7.871	4,530
PLACE1007881	5, 139 10, 863	10.414	2.393	2.603	1.012	13.782	10.374	10.918
PLACE1007885	3, 536	7.072	22.069	4.855	1.990	1.974	3.199	4.659
PLACE 1007908	63.322	28.830	20.884	16.585	18.747	22.332	15. 357	17.050
PLACE 1007922	6.729	11.816	3. 722	1.844	4.727	16.181	8.423	4.078
PLACE 1007946	27. 577	42.553	23.514	21.412	15.005	14,653	22.549	38. 209
PLACE 1007950	28. 154	21, 145	11.483	10.791	14. 345	20.195	13.857	11.448
PLACE 1007954	1. 952	1.428	1.401	0.592	0.724	0.690	1.654	- 2.786
PLACE 1007955	30.872	13.716	10.671	9. 325	3.419	15, 434	21.029	15.095
PLACE 1007956	1.554	4.401	1.470	1.778	0.511	0.943	0.995	8. 053
PLACE 1007958	23.822	7.110	10.987	1.811	8. 123	9. 545	15.981	7.219
PLACE 1007965	18.538	20.464	2.855	8.612	5. 623	10.415	21.427	13.049
PLACE 1007969	71.000	42.207	14.155	7.330	17. 492	25.314	22.985	16.519
PLACE1007971	8. 582	17.461	12. 294	9.798	9.716	7.546	12.569	20.375
PLACE1007990	14.189	22.169	6.468	9.895	22.657	6. 165	13.868	14. 027
PLACE 1008000	0.000	0.000	1.759	0.861	0.988	0.774	1.458	0.870
PLACE 1008002	0.864	4. 483	1.720	0.911	2. 225	0.000	3.225	2.113
PLACE 1008037	8.517	15. 137	4.093	2.533	2.819	5. 266	6.174 3.724	7.710 2.532
PLACE 1008044	3. 591	23.823	1.467	5. 023	1.182	19, 457		5.063
PLACE 1008045	18. 199	6.964	4. 191	3.679	17. 990	6.174 29.174	6.044 43.170	18. 085
PLACE 1008080	76.289	22.095	15.736	9.042	6.883	5.838	13.221	8. 271
PLACE 1008092	20.084	14.350	5. 254	4.007	9, 692	30.034	18.610	15.864
PLACE 1008095	66.206	18.003	15. 876 8. 653	4.784	6. 369	24.163	14.324	8.775
PLACE 1008 105	9.855	17.053	29. 282	1. 321	21. 336	190. 243	17. 482	0.000
PLACE1008107	14.915	3.951	10.948	3. 878	3.406	5,838	4. 201	5. 349
PLACE 1008111	8. 429 107. 214	70.670	30.690	73.906	24. 521	56, 386	67.918	68.831
PLACE1008113	31, 236	2.957	2. 188	2.896	3. 218	4, 599	3.943	4. 297
PLACE 1008 122	24. 832	21,510	6.892	5. 243	10.303	6.956	15.518	10.266
PLACE1008129 PLACE1008132	20. 962	34. 980	15.446	14.729	12.780	18.057	15.325	27.742
PLACE1008137	97.118	20.794	22.343	16.524	21.684	39.970	38.580	25.034
PLACE1008174	45.018	51.261	15.909	36.535	14.772	26.923	25.502	28.082
PLACE 1008177	41.484	79.290	24.754	30.372	26.003	23.816	34.010	37.711
PLACE1008181	1.719	2.220	2.731	0.000	1.579	0.000	6.557	3.286
PLACE 1008 195	59.623		14.221	11.368	19.333	17.299	34.734	21.508
PLACE1008198	30. 548		9.985	9, 568	10.838	10.004		13.967
PLACE 1008201	18. 370	7.316	4.891	5.330	6.707	8.374	16.701	15. 508
PLACE 1008209	11. 353		6.786	7.826	11, 313	9.337	6. 422	11.127
PLACE 1008226	40.512		15.314	15. 161	14. 198	15.868	18,668	19. 246
PLACE1008227	40. 507		13.616	20, 914	14.854	9.753		4.587
PLACE1008231	13.879			4. 727	8. 085 32. 492	4.880 27.046		14.304
PLACE1008238	62. 239		22.111	14. 596		4. 285		4.727
PLACE1008244	2.208				3, 829	2,643		6.790
PLACE 1008249	9. 950			14, 938 54, 336		53.728		97.747
PLACE1008266	177.598			15.300		14. 210		13.366
PLACE1008273	26.850			4. 989		2.541		
PLACE1008275				18.567				
PLACE1008280								
PLACE 1008282								
PLACE 1008297								
PLACE 1008303						4, 054		
PLACE 1008309								
PLACE 1008329								22.076
PLACE 1008329					13.083			
LEVOT 1009330	40.03	. 1						

Table 145

					30 640 1	C 771	18 146 1	10 270	14 601
	PLACE 1008331	27.986	47.595	19. 541	30.549	6,771	12. 430	19. 559	14.501
	PLACE 1008351	31, 374	25.837	26.940	15. 283	21.769	12.877	29. 581	16.720
				10.669	11.527	6. 248	8. 108	12.839	23.469
5	PLACE 1008356	11.038	24, 238						
5	PLACE1008359	23.219	18.821	4. 585	6.804	10, 499	0.000	2. 547	1.435
			12.077	7.076	8.221	6.772	3.046	4. 473	8. 994
	PLACE 1008358	7.861							
	PLACE 1008369	13.265	19.288	23. 206	5.056	9. 188	8.024	3. 252	7. 150
	PLACE 1008392	33.219	24.613	7. 199	8.079	6.094	3.416	5. 773	14. 321
					115.713	108, 082	197. 383	152.685	161.031
	PLACE 1008394	408.885	231.502						
	PLACE 1008398	25. 185	65.413	11. 186	3.178	10.520	12.052	60. 522	4. 172
10		9.122	14, 441	7. 348	5. 588	6.040	4.705	5. 467	13, 166
	PLACE 1008401						2. 926		
	PLACE 1008402	9.663	11.925	9. 911	6.799	5.684		6. 105	8.816
	PLACE 1008405	564.405	448.002	386. 959	390.811	233.214	323. 322	279. 406	299.078
				107.706	88.926	100.879	133.079	164. 162	134.635
	PLACE 1008409	310.254	194. 222						
	PLACE 1008420	102.871	44.916	30. 154	14.685	18.701	45. 968	47. 225	19.396
	PLACE 1008424	7.842	8. 421	6.860	6.448	7, 117	8.493	7. 105	5.879
15						16.885	9. 223	17. 802	15.759
	PLACE 1008426	34.481	18.699	20, 403	1.577				
	PLACE1008429	19.812	18.343	10. 368	12.697	6.738	14. 423	9. 882	12.964
	PLACE 1008430	15.959	9.694	5. 026	2.761	4, 442	8.785	16.237	9.842
		13. 333							
	PLACE 1008437	29. 520	12.626,	6.518	4.954	3, 470	6.216	6.790	9. 990
	PLACE1008453	45. 498	38. 572	11. 482	14.114	13.893	18.459	30. 671	26. 924
	PLACE 1008454	92.852	69.938	35. 812	43. 358	32.139	34. 380	44. 342	24. 973
00									
20	PLACE1008455	110.060	132.654	101.535	72.107	48.679	28. 207	49. 762	96.618
	PLACE 1008457	221.026	164.638	87.890	67.565	56.681	96.733	57. 289	64. 132
	PLACE 1008465	14.482	45. 181	6. 482	5.652	7.215	4. 989	7.987	10, 103
	PLACE 1008469	191.519	126, 151	83.503	66.767	67. 955	101.454	113.684	104.824
	PLACE 1008488	12.143	25.044	5. 332	0.377	5. 344	4. 917	4.843	10.115
			18, 134	9. 335	5.792	12. 237	15.758	18.736	15.770
	PLACE 1008519	26.949							
25	PLACE1008524	16.341	9.879	14. 963	4. 596	10.881	12.847	12.491	8. 424
	PLACE 1008531	26.300	44.215	12, 618	14.596	8.835	12.002	17.900	27,017
					12.256	5.529	20.046	14. 458	31.354
	PLACE 1008532	23.293	26.180	13, 194					
	PLACE 1008533	50.837	25.004	15.099	14.960	12.107	13.885	15. 331	16.687
	PLACE 1008542	7.209	11.351	11.148	11, 159	7,406	3. 275	4.870	8.793
						35.339	4.849	21.994	15.899
	PLACE 1008549	24.848	27.469	14. 722	7.446				
30	PLACE 1008560	16. 248	9.601	10.580	4. 328	6.786	9.843	14.007	6.753
30	PLACE 1008567	31.376	45.822	16.034	16.944	14.791	13.929	17.148	17.570
						15.738	22.714	15. 252	13.545
	PLACE 1008558	9.263	28.507	12. 536	7.903				
	PLACE 1008569	21.434	13.045	5.050	6. 520	8.664	9. 142	10.799	9.664
	PLACE 1008584	29.627	24.002	13.657	10.990	11, 106	13.734	22.655	21.057
					7.124	8. 320	13.100	8. 184	14.617
	PLACE 1008585	25.861	23.246	13.959					
2=	PLACE 1008603	11.593	12.897	3.634	5. 109	4.753	7.887	18.167	11,774
<i>35</i> ·	PLACE 1008621	6.723	3.752	3.073	2.882	0.628	2.394	2.356	6.079
						1.816	1.254	2.598	3.068
	PLACE1008625	5. 997	8.406	1.768	1.055				
	PLACE1008526	5. 484	3.562	1.402	1.123	1.403	3.049	5.665	3.510
	PLACE 1008627	49.718	18.742	10.960	7.037	8, 831	13.117	21.039	15.675
	PLACE 1008629		10.172			9, 612	12.027	17.865	12.171
		21.102	28.942	11.982	3.365				
	PLACE 1008630	9.527	21.990	10.098	9.473	7. 038	5. 568	7.548	9.704
40	PLACE 1008643	41.545	29.478	16. 220	15.566	9. 566	16.636	24.733	18.160
					2.532	0.932	2.348	3.778	2,601
	PLACE 1008650	4. 202	2.471	1.051					
	PLACE 1008657	10.667	15.060	5. 999	8.523	5.606	4. 350	8.873	8. 539
	PLACE 1008564	7.147	9.457	8.348	2.448	3.877	5.707	7.490	2.436
	PLACE1008693	35.830	32.008	13. 154	7.301	10.960	12.214	13.885	10.914
	PLACE1008696	30.598	14. 195	9.900	6.913	8.747	8. 454	9.419	10.479
45	PLACE1008715	6.265	13.318	2.170	5, 131	3,050	3.374	6. 120	5.989
	PLACE1008718	10.756	11.071	14. 349	7, 225	9,919	5. 434	16.844	11.965
			11.0/1			+ 3, 313			
	PLACE1008722	19, 150	29.145	12.082	14, 107	7.317	7.365	11.291	13. 128
	PLACE1008738	12.649	24.539	11.238	5.658	9. 182	17. 327	16. 429	12.149
							8.825	6.516	10, 105
	PLACE 1008742	4.334	14.217			5. 946			
	PLACE 1008744	8.130	10.071	2.674	2.854	2. 153	2.940	3.519	4.369
50	PLACE 1008748	8, 135			1.850	7. 331	2.772	2.033	6.870
50			6.332						
	PLACE 1008757	0.000	1.927	1. 248	0.983	2.427	2.818	1. 135	1.993
	PLACE 1008766	4.606	24.202	3. 522	1.672	4.576	4.758	5. 053	2.617
	PLACE 1008785	84.472	51.726	24. 136	25.096	17. 140	24.917	15. 172	19.772
	PLACE 1008790	31.403	25. 252	14, 095	12.995	13.157	12.786	21, 229	14.549
	PLACE 1008798	3.470					1.268	3.684	4.700
E E	LEVER 1000139	1 3.410	1.133	1 6. 113	1.677	1 2.007	1.200	7.004	1 ,,,,,,,
55									

Table 146

			0.000	- 212 [	3 300 1	4 163 1	C 30C 1	0.064	7.993
	PLACE 1008807	11.746	9.388	7.010	3, 398	4. 152	5. 285	9. 954	
	PLACE 1008808	10. 497	2.010	1.832	1.724	2.154	0.000	2. 960	3. 938
	PLACE 1008813	43, 335	8, 124	3, 170	3, 472	6.648	0.000	5. 265	5. 081
5		13. 208	30. 377	8.014	11.550	8.774	9,700	18. 296	15. 156
3	PLACE 1008836				38.594	17.658	12. 467	22.869	17.982
	PLACE 1008851	35. 131	44.912	12. 581					8.631
	PLACE 1008854	5.882	9. 135	0.000	4.861	5. 302	11.675	6. 631	
	PLACE 1008864	48.984	42, 179	18.396	30. 397	21.064	16.595	22. 139	23.902
	PLACE 1008867	12.377	56.824	11.324	9. 452	18.933	14.620	10. 186	16.826
		49.946	97. 258	18. 984	54.608	14.811	25. 438	23. 529	37, 995
10	PLACE 1008876			15, 208	16.042	20.811	10, 479	15. 115	15. 164
10	PLACE 1008887	26.489	38.089			19.625	2.141	5.762	6.510
	PLACE 1008902	22.685	13.678	2. 921	7. 383			11.666	13. 828
	PLACE 1008911	9.060	33.193	12. 197	13.855	16. 972	6.042		
	PLACE 1008917	42.217	35. 405	16.607	7. 150	18.874	11.592	41.024	22.806
	PLACE 1008920	32, 162	3. 225	1.754	3.766	3. 590	9.067	6.073	2. 425
		13.417	17.966	5, 400	5. 416	6.761	6.566	12. 223	7.803
4.5	PLACE 1008925			9. 408	9. 552	6,095	6. 477	15.830	11.057
15	PLACE 1008930	15.886	28. 504				6, 117	11, 146	9. 917
	PLACE 1008934	23.759	18.548	12.356	6. 943	12.662			
	PLACE 1008941	8, 316	9.677	5. 776	9. 338	9. 104	5. 758	12.723	13.555
	PLACE 1008947	150.057	83.432	44, 128	33.278	56.786	59.699	86.640	63.955
	PLACE 1008984	8.712	10.873	4.711	5. 382	2.608	4.656	10.459	7. 929
		25. 866	40. 327	13.608	7.899	8.177	11.454	23. 995	16.883
	PLACE 1008985				2. 403	2.775	3.796	8. 332	3.014
20	PLACE 1008994	18. 162	8. 786	5,711			6. 161	11. 355	7.439
	PLACE 1009020	11.578	10.784	5. 965	4.614	3.880			
	PLACE 1009027	21.125	15. 947	4.623	2. 459	3.520	11.909	6.684	4. 839
	PLACE 1009039	8.664	10.154	6, 735	2. 521	7.750	11.874	23.006	4. 885
	PLACE 1009045	23.977	20.675	6, 979	7.407	4.810	5.799	35. 292	9. 408
		5, 091	10.171	2.268	5. 954	4, 362	0.000	5.318	6. 521
	PLACE 1009048				4.708	3.880	0.000	4. 164	8. 669
25	PLACE 1009050	3. 470	5. 590	9.098				23. 124	21.665
	PLACE 1009060	34. 280	32.398	9.016	17.645	9.108	20.791		
	PLACE 1009067	55.833	32. 552	13.821	3.577	11.693	36.606	50.944	44. 507
	PLACE 1009071	137, 113	72.622	42.839	42. 259	33. 328	32.445	60.967	59.816
	PLACE 1009090	30.957	25.567	12.139	8, 147	11.883	22.624	22. 381	10. 572
		42.486	15.715	10, 526	6. 902	14, 110	5. 159	15, 660	17. 580
	PLACE 1009091				8.271	10.714	15.361	21.919	17.604
<i>30</i>	PLACE1009094	21.335	70.138	13.676		8. 281	12.542	10.801	31.093
	PLACE 1009099	7.525	13.610	8. 280	12.776				
	PLACE 1009110	13.415	6.006	4. 409	1.648	2.849	4.580	4. 965	5. 369
	PLACE1009111	67.629	16.954	11.182	7.515	0.000	7.804	15. 142	12. 395
	PLACE 1009113	10,615	8.546	4, 331	4.640	5. 385	6.432	5. 643	10. 147
	PLACE 1009130	6.901	19.609	23.895	6.666	2.762	2.544	1.446	3. 744
			20.426	5. 736	7.683	3.673	7.990	4, 988	8. 429
35	PLACE1009150	13.031			55. 149	41.987	39.328	50.150	57.022
	PLACE 1009155	72.157	61.300	57.610		11.626	14.318	26. 507	16. 570
	PLACE 1009158	28.497	16.235	13. 335	10. 201			27. 587	24.065
	PLACE 1009166	58.030	29.706	24. 997	22. 721	18.028	18. 384		
	PLACE 1009172	16. 222	19.005	7. 161	4. 843	6. 408	6.734	8. 370	6.017
	PLACE 1009174	50.892	48.998	32.343	28. 578	23. 381	21.627	24. 363	21.250
	PLACE 1009183	61.545	60.739	14.751	35.658	16.796	15. 529	13.831	15. 373
40	PLACE 1009186	5.029	11.552	6. 154	4. 372	2.812	8.067	6. 125	4. 542
				2.215	1.077	0. 922	0.000	0,000	0.879
	PLACE 1009190	0.112	1.383			5, 301	4.199	8, 229	7. 836
	PLACE 1009195	15.938	15.069	6.337	11.235				29. 814
	PLACE 1009200	56.062	49. 582	26.621	32.612	20.016	13.451	19. 592	
	PLACE 1009217	9.045	7.250	3. 382	10. 839	3.645	4.052	6. 924	17. 092
	PLACE 1009230	35.137	34.356	13, 699	21.015	16.141	8.394	19.789	7.528
45	PLACE 1009236	34.867	17.528	8. 326		9.004	12.493	27.327	9.172
	PLACETOUSZS			28, 320			9.019	29, 697	24. 935
	PLACETO09246	51.787	71.164				43.325	58.877	30.908
	PLACE 1009265	92.450	36.053	21.026					
	PLACE1009279	25.174	8. 294	11.814			10. 155	13. 253	6. 328
	PLACE 1009298	28.708	18.088	16.943	10.646		14. 708	8.886	9. 738
	PLACE1009308	175.031	34.217	34.842		32.150		72. 179	28. 297
50	PLACE 1009319	21, 209	35.386	7.874			12. 353	8.009	11.881
							16.474	11.081	13.533
	PLACE1009328	34.584	30.370	22.052				4. 162	27.779
	PLACE 1009335	3.869		12.941					
	PLACE 1009338	4.629	13.280		2.945	5. 427		7. 332	6.665
	PLACE 1009344	33.854	26.440	7.150	7.043	5. 231		17.883	
	PLACE 1009355	10.104					6. 227	6.669	19.235
55	[ CAGE 1003000	1 .0.,04							
55									

Table 147

					- A A S S - I	-7 101 T	7 006	16 CAC 1	0 444
	PLACE 1009368	42.051	14.861	10.631	6. 209	7.101	7.025	15. 596	9.443
	PLACE 1009375	19.461	10.862	1.937	2.151	5. 975	8.807	9.665	4.779
	PLACE 1009388	41.922	22.694	9, 119	6.828	8.777	12. 117	12.174	12.815
5	PLACE1009398	9,410	16, 113	10.077	14, 136	8, 930	7.363	10.053	24. 623
J		27. 332	38221	. 8. 577	12.742	8.050	16.604	26.279	8.093
	PLACE 1009404		6, 807	2.954	3.849	2, 292	2.641	6. 326	4. 231
	PLACE 1009410	9.672			7.745	9. 842	13, 265	12.016	15, 150
	PLACE1009417	11.321	13. 342	11.760			145.780	98.718	82. 459
	PLACE1009424	143.874	161.949	83.678	44. 296	55. 295			
	PLACE 1009434	3.639	9.793	2. 953	4.133	3. 385	3.038	6.041	7.934
10	PLACE 1009443	10.126	5.900	2.564	1,418	3.826	4. 205	6.190	3, 051
	PLACE 1009444	75.456	51.672	32.690	29. 162	27.896	35, 657	35.401	25.671
	PLACE1009459	110.550	32, 136	23. 433	13.124	19.500	46.330	49.514	27. 422
	PLACE 1009460	7.804	18, 196	5. 042	1.388	3.715	11.334	3.840	5. 965
	PLACE 1009468	24.940	28, 488	14.998	8.351	9, 763	9. 596	31.733	20.845
			12.973	6, 635	5.700	3. 950	6.971	12.745	5. 157
	PLACE 1009476	18. 955			21.520	9. 248	17.462	14.475	15. 028
15	PLACE1009477	28. 528	28.026	14.306		1.914	6.774	6. 193	9. 481
	PLACE 1009493	9.706	13.481	2.399	3.953				
	PLACE 1009502	3.768	2.155	2. 938	0.891	2. 156	5.093	2. 120	1. 962
	PLACE 1009524	41.369	7.099	18.781	3.777	7. 184	16.229	19.248	12. 391
	PLACE 1009527	41.383	14.310	8.219	3.634	8.710	15.448	19. 901	11, 203
	PLACE 1009531	43. 331	29, 448	11.293	13.089	12.741	23.938	25. 244	25. 592
	PLACE 1009535	11.347	16.999	7.257	9, 551	6.031	5.821	7.459	13.160
20	PLACE 1009539	27.355	33. 924	17.760	19.107	12.625	17.181	18. 261	21, 706
	PLACE 1009540	26.063	18, 180	18, 706	13.776	10.936	19.307	24.429	16.284
		32.314	9. 517	6.333	8, 159	7.348	18.062	22. 235	9. 384
	PLACE 1009542			5.625	1.298	3. 320	4.724	8. 207	4.406
	PLACE 1009546	12.399	7.380			6. 465	10.966	16. 358	19. 196
	PLACE 1009556	13.954	15.082	5. 948	1. 391		8.233	9. 438	12.507
25	PLACE 1009569	22.909	21.209	6.670	12.434	5.803			
25	PLACE 1009571	13. 458	10.535	6.868	4.758	5. 027	9.733	7. 553	9. 107
	PLACE 1009573	16. 235	9.693	6.699	13.447	6.873	4.277	8. 380	12.992
	PLACE 1009576	4, 851	10.697	8. 157	4.542	2.949	3.677	5. 201	5. 143
	PLACE1009580	35. 237	47.578	24.938	26.636	15.366	25. 243	27.920	23, 541
	PLACE 1009581	30.483	8.604	7.654	6.565	7.711	16.692	24.706	13. 168
	PLACE 1009587	3.476	3.868	5.230	3.387	4.099	4.838	6.514	5.783
30	PLACE 1009593	7. 424	8.043	3, 949	5.143	4.859	7.848	5.031	4. 525
	PLACE 1009595	63.588	58.749	27.289	26.946	25. 118	25. 486	32.674	29, 915
	PLACE 1009598	10, 136	8.803	2.554	6.077	10.559	3.608	12.421	11.189
	PLACE 1009 600	15. 391	21.884	10.853	7.573	11.964	20.158	11. 161	14.987
		32. 270	9.947	13.494	11.363	10.658	9, 443	19. 197	18.000
	PLACE 1009604				26. 439	26. 445	29.568	26. 158	30.122
	PLACE 1009607	75. 364	85. 156	35.035			2.792	2.408	6.068
35	PLACE 1009613	4. 353	6.164	2.640	5, 243	1.911		22. 997	15. 101
	PLACE 1009621	29.001	49.946	14, 693	13.116	18. 138	23.193		
	PLACE 1009622	27. 300	10.327	8. 159	5.651	12.385	9.234	15. 408	7.132
	PLACE 1009624	27.426	19, 103	3, 360	2.878	7. 125	4, 125	12. 179	7.539
-	PLACE 1009637	5.028	13.109	5.041	2.366	9.802	4.190	6.416	4.450
	PLACE 1009639	9, 956	16. 237	4.056	3.880	8.587	3.660	14.640	27.577
40	PLACE 1009654	29.616	69.766	58.647	5.371	50.183	22. 307	21.782	12.466
40	PLACE 1009689	10.143	12.022	13.185	10.544	15. 157	2.663	7.467	4.763
	PLACE 1009665	19.662	15,718	10, 263	8.654	15.968	3.947	7.286	5.058
	PLACE1009569	74. 335	65.299	22.539	17.665	23.035	35.889	47, 853	26.094
	PLACE 1009670	48.759	30, 681	15. 505	15, 680	13.512	21.863	46.277	13.806
		9. 584	14.533	5. 232	5. 640	7. 390	7. 392	11.586	7.014
	PLACE 1009708				1.030	1. 425	4.841	5.611	3.780
45	PLACE1009721	0.000	5.965	1. 997	13.383	9. 274	42.308	14, 822	16.604
	PLACE 1009731	31.531	29.697	5. 222	13.303		11.001	10.728	17.147
	PLACE 1009735	24.842	17, 444	8. 225		3.900			
	PLACE 1009737	20, 121	19.390			4. 987	10.582	13.461	11.206
	PLACE 1009741	3.834	48. 256	3.058		12.402	22.656	1.749	4.187
	PLACE 1009752	37. 588	360.319	9, 532	24. 594	5. 279	91.807	22.992	435. 143
	PLACE 1009763	15. 243				11.844	8.197	7. 432	17.382
50	PLACE 1009766	15. 481				8.733	9.416	11.841	13.177
	PLACE 1009772	25. 177				5.178	8.892	12. 233	6.915
	PLACE 1009782					13.633	6.484	6, 993	16, 851
	PLACE 1009794							17, 171	7.455
	LEVOS 1003134							6. 535	
	PLACE 1009798	16. 321	14, 039		11.317				7. 202

Table 148

			4 333	2.997	5. 329	2. 393	2.613	22.333	11.323
	PLACE 1009845	15.220	4. 333	17, 197	64.071	16. 467	30.251	38. 997	341, 202
	PLACE1009849	44.946	194, 619		9,010	7.686	17.560	23, 505	10, 157
	PLACE 1009857	21.842	11.784	14.813	38, 531	21.999	12.526	21. 181	42.147
5	PLACE 1009861	55.060	52. 334	22.982		19.845	26, 217	21.062	57. 158
	PLACE 1009872	42.867	65. 398	11.814	72. 397			20.382	36. 147
	PLACE1009877	144, 154	73.771	52.613	35. 986	26.345	12. 461	20. 282	9. 265
	PLACE 1009879	31.357	19. 333	43.105	15.026	16.781	19. 583		1.893
	PLACE 1009886	3.579	8. 567	2.869	1.043	1.021	1.571	2.025	6, 285
	PLACE1009888	10.362	6. 906	3.541	1.720	7. 325	6.831	7.680	
10	PLACE 1009908	16.750	13.979	9. 123	5.093	6.107	7.524	13. 900	10.848
	PLACE 1009919	25, 958	16.368	12.802	7.838	12.682	8.032	7. 157	13.099
	PLACE 1009921	5, 294	5. 301	2.647	2. 379	6.669	1.694	6.864	1.626
	PLACE1009923	7,666	10.700	2.427	3. 962	7.335	13.971	11.821	6.627
	PLACE 1009924	26.023	5. 583	3.961	1,712	2.571	0.000	6.021	12.826
	DI 100 1003 324	3, 609	1. 404	0.882	1.882	0.508	7.012	3. 169	2. 473
4.5	PLACE 1009925	37.980	53.080	21.843	35.590	14.645	26, 179	18. 163	39.695
15	PLACE 1009931		3. 468	2.666	1.324	0.764	2.382	4. 922	3.501
	PLACE 1009935	7.854	21.773	17.615	11.373	13. 359	12.852	18.329	12.383
	PLACE 1009947	44.482		3.780	2.246	7. 199	6,513	3. 962	22.636
	PLACE 1009961	3. 264	4. 537,		9.558	7,736	13.999	13.695	8, 124
	PLACE 1009971	24. 201	14.113	8.964	17. 226	11.857	37.703	32.523	24. 126
	PLACE 1009982	90. 204	37.402	17.490		5. 980	18. 389	18. 358	10. 327
20	PLACE 1009992	32.659	8.657	9. 454	6.512	15.829	28, 418	28. 296	25. 865
	PLACE 1009995	21.779	25. 489	20.929	6.918	12.955	27.052	15. 574	19.395
	PLACE 1009997	39.778	25.957	22. 163	18.804			6.047	4.825
	PLACE 1010002	7.208	6.675	2.154	3. 335	4,711	2.649	16. 561	3, 132
	PLACE 1010011	15, 700	11.002	2.148	0.691	4. 571	3.619		9.751
	PLACE1010013	18.169	7. 231	5. 446	11. 205	1.374	6.028	15.057	
05	PLACE1010021	9.423	11.541	8.788	5. 901	5.744	6. 434	11.142	6.794
25	PLACE1010023	48.546	20.475	6.683	8. 439	7.872	6.849	14, 748	19.147
	PLACE1010031	23. 253	23.746	12.677	11.119	9. 178	23.991	11. 578	15.444
	PLACE 1010039	8.216	5. 363	3.410	2.754	3.443	3.809	2. 994	3.074
	PLACE 1010045	28.520	20. 935	14. 936	23.387	11.939	11.927	10. 256	27.268
	PLACE 10 10 053	11.420	12.399	2.211	6.506	4. 422	6.813	4. 552	6.626
	PLACE 1010060	61.784	35. 230	25.530	15, 116	15.866	30.074	32.753	19.303
30	PLACE 1010069	13.551	3.560	5, 924	2,419	1.178	3.632	7.745	6.202
	PLACE 1010070	12. 192	12.514	5.728	3.839	6.386	6.674	3. 922	9.645
	PLACE 10 10074	58. 736	80. 938	44.955	39.497	35.506	33.481	44.710	58.097
		241.223	62.057	77.062	19.863	\$9.519	134.094	156.661	51.913
	PLACE 1010076	85.849	25. 937	22.479	14. 142	12.854	44.885	40.845	17,940
	PLACE 1010078		6. 981	0.000		0.000	0.000	11.595	5.593
35	PLACE 10 10 08 1	0.000	20.600	7.478		3, 550	9. 386	8.886	8.608
00	PLACE 10 10083	27. 240		5. 452		8.073	1.380	4. 234	6.582
	PLACE 1010089	10.050				6.620	5.621	14.446	21.698
	PLACE 1010096	15.851	23. 598	8. 484		10.558	11.564	11.290	13.860
	PLACE 1010102	15. 331	13. 251	7.699		17.745	14. 276	23. 241	18.148
	PLACE 1010105	35. 995				19. 261	13.790	13.444	18.632
	PLACE1010106	22.316		22.970			19.911	32.607	24.027
40	PLACE 1010130	31.537				10.772		10. 934	9. 482
	PLACE1010132	29. 236				5. 570	12.883	17, 154	
	PLACE 1010134	33. 947				7.730	13.218	448. 388	
	PLACE 1010139	598.413				206. 627	352.839		
	PLACE 1010148	3, 132				4. 356	4. 931	4. 453	
	PLACE 1010152	26. 445					7.409	13.853	
45	PLACE 1010155	373.743					17.546	25. 269	
-	PLACE 1010156	9.490		7.14					
	PLACE 1010161			2.530	5. 962				
	PLACE 1010181	5, 29				6.824			
	PLACE 1010194								
	PLACE 1010202							19. 48	
								5.65	7.547
50	PLACE 1010231								3.884
	PLACE 1010235								
	PLACE 1010237								
	PLACE 1010251								
	PLACE1010261								
	PLACE 1010270	3.52	8 2.56	- 1 4.68	7 1.01	- 6. 911	3.00	. ,	

Table 149

			10 200 I	E 456 T	7.563	10.408	11.696	11.805	8.650
	PLACE 10 10273	18.198	10.799	5.456		13.997	14, 739	14. 496	20.193
	PLACE 1010274	20. 202	18. 193	10.486	9. 941				10.102
	PLACE 1010277	8.937	117. 446	5. 398	3. 512	4.011	1.815	8. 164	
5	PLACE1010293	60, 036	62. 516	18.939	20.260	11.120	8.879	14.863	20. 261
	PLACE 1010297	10, 456	6. 185	4.720	3.674	4.733	7.175	13.007	16.488
		17.008	24. 187	7.187	17, 501	12.198	7.801	13.650	16.733
	PLACE 1010300		200. 863	167.599	97.554	142.759	225.854	230, 002	179.252
	PLACE 1010310	413.605		16.701	11.196	12.716	16, 372	17. 343	18.872
	PLACE1010321	36.500	66.804			3.447	3.169	4, 956	2.116
	PLACE 1010324	0.000	8. 637	3.654	1.998	11.246	12.390	12.923	13.038
10	PLACE 1010329	30.906	39. 387	9.407	14.862			103.997	33.527
	PLACE 1010330	182.247	52. 788	42.842	20. 272	46.381	81.215		
	PLACE1010335	20.429	27.007	19. 301	14.056	19.661	14.766	38. 945	15.803
	PLACE 1010341	15.512	16.397	5. 370	6. 587	7.213	4. 477	8. 264	8. 294
	PLACE 1010342	5.485	5.818	1.359	2.968	2.444	1.600	5.664	2.119
	PLACE 1010346	27.509	31, 551	11.234	16.701	10.972	11.936	17.856	16.679
			42.341	20. 410	19.994	20.516	14.039	23.979	21.826
15	PLACE 1010362	37.800		4.890	1.792	3.344	6.491	9.839	7.931
	PLACE 1010364	3.637	7.688			60.226	64.663	105.004	71.078
	PLACE 1010368	160.448	94. 255	69.658	48. 395	19. 197	22, 390	35.216	29.763
	PLACE 1010373	50.531	36.656,	15. 978	12.876		11.968	26. 253	22.869
	PLACE 1010383	60.222	42.672	28, 248	34.317	29.853			0.000
	PLACE 1010385	0.000	3.211	0.000	1.653	2.697	0.000	3. 102	
	PLACE1010389	45.010	32.965	23.673	18.734	15. 387	31.864	30. 482	22.113
20	PLACE 1010401	12.554	12.082	7. 358	3.809	4.486	9.049	11.163	8.023
	PLACE 1010410	46.622	19.531	23.525	15.038	12.094	26.576	30, 580	20.641
	PLACE 1010418	63.170	54. 185	47. 245	53.690	29, 885	39, 952	35. 428	40.754
•		8, 496	10.271	8.511	8.469	6.845	60.372	16.883	14.750
	PLACE 1010425			76, 495	49.901	31.535	91.673	163.084	100.340
	PLACE 1010443	139.820	68.717		41.679	24. 598	29.543	39.397	42, 435
25	PLACE 1010445_	55. 230	63.853	40. 195		7.811	11.242	18.708	11.397
23	PLACE 1010481	25.071	14.236	12. 932	6. 994	7, 434	27.561	32.378	14. 322
	PLACE 1010482	62.044	30, 485	12.054	12.510			8.897	7.016
	PLACE 1010491	6.692	11.769	7.835	7. 107	2.403	7.772	11.758	
	PLACE1010492	8.815	25. 244	14. 396	11.795	10.757	10.883		14.039
	PLACE 1010509	8.728	8.603	9.041	7.620	3.097	8. 537	33. 229	11.438
	PLACE1010518	53.737	47.379	37.510	43.651	35. 422	32. 152	29, 681	41.839
30	PLACE 1010522	74,460	121.326	35, 701	24.026	30.767	37. 996	82.263	44.005
••	PLACE1010529	13.116	47.273	16.874		11.833	10.805	15.047	19.475
			15.015	13.620		6.861	9,050	12.611	9, 113
	PLACE 1010547	10.791		20. 254		7.397	21.958	19.638	14.752
	PLACE 1010560	36.084	24.074	13.412		8.160	11.786	18,067	7.693
	PLACE 10 10562	21.600	13.040				6.175	8, 453	8.370
	PLACE 1010579	5.809	7.166			14. 505	33.521	41.838	28.526
35	PLACE 10 10580	50.738	35. 579				12.163	6. 932	13.134
	PLACE 1010599	22.697	6.399				8.754	6.341	10.710
	PLACE 10 10606	17.463	9. 467			13. 966			17.679
	PLACE 10 106 16	16.337	36.535			9.645	7.170	14.986	
	PLACE 1010622	30.437	14.238		5.708	14.881	22.701	23.807	13.849
	PLACE 1010624	25. B23	18.627	12.823				11.721	14.514
40	PLACE1010628	13.901	8.075		7.978				11.287
40	PLACE 1010629	27.634							10.210
	PLACE1010830	12.405							
		19.768							
	PLACE 1010631	61. 423					34.204	25.544	13.578
	PLACE 10 10651								
	PLACE 1010661	34, 409							
45	PLACE 1010662	25.892							
45	PLACE 1010668								
	PLACE 1010702	18.288							
	PLACE 1010709		137.91						
	PLACE 1010713			5 14.08					
	PLACE 10 10714				1 3.91	2 6.92			
								12.608	13.906
50	PLACE 10 10716								
	PLACE1010717								
	PLACE 10 10720								
	PLACE 10 1073								
	PLACE 101074	9.10	1 4.61	0 3.58					
	PLACE101075					7 15. 19	8 31.01	0 28.79	3   18.098
	1. CACC101010								

Table 150

				45 444	07 70E 1	12 691	36 360 1	19. 393	56.313
	PLACE 1010761	26. 459	111.645	25. 313	97. 785	46. 971	35. 398	41. 271	23. 174
	PLACE1010771	97.575	46.358	30.540	20. 492	21.112	45.643		12.720
	PLACE 1010784	34. 813	13.196	12.948	5. 263	8. 395	17. 778	16. 235	30.768
5	PLACE 1010786	35. 506	55. 424	19.835	19, 203	16.991	22. 191	24.116	
	PLACE1010789	14.662	9.740 1	10.856	8. 035	6.035	6.662	6.785	6. 617
	PLACE 1010800	12.898	11, 478	12.969	11.574	8. 280	13.756	9.074	10.785
	PLACE 1010802	9. 976	7. 839	11. 257	6.385	8.708	6.482	9.517	9.615
	PLACE 1010811	6. 267	10,750	6.130	5. 326	2.131	5, 807	7. 023	5. 806
	PLACE 1010813	107.134	54. 846	41.785	19.939	26.019	51. 877	79.848	45. 993
10	PLACE 1010827	11.543	12.554	6.090	2.687	4.360	10.117	10.344	9.099
. •	PLACE 10 10833	70.712	36. 952	36.612	16.799	28.163	60. 904	40.462	31.469
		56. 261	52, 196	32.723	40. 363	32.757	24.743	30.658	33.056
	PLACE1010839	15, 444	56. 200	14.751	17.041	11. 951	20.702	14.170	62.029
	PLACE 1010856		24.574	22.222	15.965	8.058	13.468	10.994	20.619
	PLACE1010857	16. 284	15.311	10.708	17.750	6.704	9, 120	10.270	16, 911
	PLACE1010870	11.360		12.897	9.474	11.687	13.857	6.866	12.944
15	PLACE1010877	12. 253	23. 451	15, 696	9, 810	8. 334	17.859	26.634	77.062
	PLACE 1010882	24. 453	43.270		7.840	6.799	5. 426	7, 441	5. 870
	PLACE 1010891	12.636	7.098	6.674	16. 507	18.760	17. 466	22.357	29, 192
	PLACE 1010896	35. 110	39.870	19. 987		25. 080	27. 551	37. 245	35. 556
	PLACE 1010900	50.692	63.882	25. 595	31.970		13. 485	21.811	16.868
	PLACE 1010916	17. 218	31.574	12.713	10.089	10.861	5. 344	6. 106	5. 656
20	PLACE 1010917	8.779	3.044	3. 185	15.098	6. 120		23. 409	11.594
	PLACE 10 10924	25.229	20.092	9. 911	8.013	6.493	10.958	17.175	29. 154
	PLACE 1010925	49.823	61.948	23. 489	34. 123	17.969	19. 262	29. 924	33.816
	PLACE 1010926	49.767	50.605	22. 959	20.111	18.009	24.065		
	PLACE 1010942	85.218	46.665	26.680	26.313	22.818	28.713	30. 538	39.140
	PLACE 1010943	316.403	113.988	93. 186	72.867	91.388	149.579	188, 191	112.743
05	PLACE1010944	48. 129	50.381	15. 305	17.574	14.904	18.649	33.779	24. 850
25	PLACE 1010947	51.058	49.164	23.114	19.450	15.597	21. 983	21.814	17. 333
	PLACE 1010954	73.590	77.560	34.775	41.312	25.097	30.688	27.071	36.359
	PLACE1010960	5. 163	5.378	16.789	7.998	6.612	8.441	8. 411	7.942
	PLACE1010965	12.476	21.628	7.886	8. 825	4. 194	19.265	13.526	8.153
	PLACE1010968	34.696	21.848	9. 662	5. 337	11.298	19.848	21.002	15. 864
	PLACE1010978	34.271	21.883	15.077	11.695	13.575	20.670	28.798	23. 174
30	PLACE1010982	11.927	20.104	5. 539	9. 523	4, 555	9, 333	27. 370	20.028
	PLACE 1010990	23.709	22.125	15.859	11.150	14. 185	15.589	24.495	18.421
	PLACE 1011017	14.795	20.170	18.473	19.079	18.837	31.530	20.694	25.609
	PLACE 1011019	60.412	29.348	19.512	15.616	21.011	29.657	32.510	15.026
	PLACE 101 1026	6.403	27.542	4.006	7.156	5. 587	9. 352	6.378	23.067
	PLACE 101 1032	22.416	44.013	12.767	14. 147	7.488	10, 513	12.024	9.185
35	PLACE 1011041	43.649	29.675	20.339	13.342	17.790	18.671	26.478	21.550
	PLACE 101 1045	48.770	37.661	20.984	15.020	24.758	23.731	42.534	24.019
	PLACE 101 1046	49.343	48. 382	29.451	17.863	35.583	26.848	35.241	25.655
	PLACE 101 1054	107.000	92.094	47. 988	57.849	58.878	38.779	50.411	53.030
	PLACE 101 1056	226. 902	159.857	111.396	119.852	115.390	99.976	141.062	137.522
	PLACE 101 1057	5. 333	7. 254	4.880	6.072	5. 943	6.298	5.741	6.082
40	PLACE 101 1059	9. 231	13.844	6. 945	5. 804	7. 325	8.493	13, 139	9.998
40	PLACE 101 1066	24. 382	54, 196	22.706	25. 109	25. 646	15.697	16.286	16.716
	PLACE 101 1087	58.783	144.018	41.548	46.968	28.518	50.611	45. 100	50.864
	PLACE 101 1090	53.056	143, 896	45. 260	34.467	50, 933	95.133	280.440	58. 429
	PLACE 101 1109	75.794	119.843	42.881	49.952	43.765	33.319	35.583	34. 429
	PLACE IOI 1114	65.656	71.805	22.254	8.641	15,726	26.074	50.404	27.034
		145. 171	37. 399	52. 539	10.533	21.813	95.905	74.823	26, 509
45	PLACE 101 1116	18, 160		14. 154		7, 536	12.531	122.844	13.983
	BL - APIA	34. 682						23.747	
	PLACEIUIII33							61.612	
	PLACE1011134	63.554						18.023	11.713
	PLACE1011143	25.496						75.016	44, 532
	PLACEIDI1146	137.473							
50	PLACE1011160	24.414						28.311	19.373
30	PLACE 1011165	34.715						18.184	13.882
	PLACE1011181	50.804		25. 933	11.931	14.943		29.663	
	PLACE 1011185	98. 259					45. 963	38.543	
	PLACE 1011186	40. 892					28. 187	25.736	15.462
	PLACE 101 1203	3. 103						1.948	1.730
	[FEACETOT 1203	3. 303	2.301	7.000					

560

Table 151

							16 100 1		22 616
	PLACE1011214	19.000	30. 499	15. 354	20.715	13.540	15. 163	22. 508	23.615
	PLACE1011219	50.422	59.474	25.989	22.358	10. 192	25.888	34.747	35. 987
	PLACE1011221	13.282	16.503	9.149	13.164	7.054	10.970	12. 103	12. 291
5	PLACE1011229	21, 300	24.016	24.142	11.920	8.874	11.425	16, 577	18.885
3			22.558	21.088	13.366	17.790	47.373	24. 485	19.912
	PLACE 1011231	57.691							74.012
	PLACE1011236	145.860	58.617	57.365	30.780	34. 641	68. 303	110.808	
	PLACE1011247	65.406	45. 970	27.363	22.989	18. 925	54. 590	38. 380	66.200
	PLACE 1011263	18.980	16.439	15. 299	13.023	14. 184	8. 485	11.883	15. 956
	PLACE 1011273	3, 117	3, 517	3.011	3.406	3.973	1.889	2.488	3.416
10	PLACE 1011278	99. 532	58, 735	53. 312	77.774	36.414	56.820	55. 573	49. 298
70		65, 724	17. 465	19, 765	15. 982	16.860	28.472	40. 138	23.783
	PLACE 1011289				18. 526	59. 460	122.088	150. 314	35.889
	PLACE 1011291	162.344	63, 584	63.268			31.931	32. 378	
	PLACE 1011296	60. 289	35. 108	32.914	21.911	20. 435			27. 683
	PLACE 1011310	12.375	27. 199	12.116	10. 122	5.617	13.629	12.674	18.490
	PLACE 1011311	31.445	29. 424	19.821	36. 262	15. 558	31.421	31.132	47.294
15	PLACE 1011321	48.851	39. 888	19, 447	21.568	15. 965	16.409	16.955	23.945
	PLACE 1011325	25, 927	17.098	14.860	7.351	9.021	14.507	18. 423	13.442
	PLACE 1011332	7. 973	13. 581	8.965	7, 176	12.435	8. 470	9. 437	8.966
				94. 222	121.189	70.843	83. 242	78.735	123, 304
	PLACE1011340	135. 172	94. 377,			20.697	18. 407	11.957	19.750
	PLACE1011353	20. 244	35. 898	18. 659	17.306				
	PLACE1011360	36.650	86. 365	30. 582	12. 233	27. 406	21.790	20.034	18.573
20	PLACE 1011364	63.297	27.430	46.019	13.619	40.083	41.537	38.082	14.810
	PLACE 1011365	14. 275	15.778	11.893	10.572	8. 300	11.140 -	12.421	11.796
	PLACE 1011371	101.501	43. 555	36.081	20. 272	24.602	47.751	80.751	51.543
	PLACE 1011375	11.873	15. 442	10.915	7.912	7.069	8. 106	9.387	8.664
	PLACE1011386	207.095	98.628	73.492	39.642	58. 242	102.702	134.598	72.968
	PLACE 1011399	12.717	15.843	7.670	9.849	6.144	8.677	7. 308	6. 939
	PLACE 1011406	60.080	56. 205	37. 483	22.859	14.794	35.277	39.952	36.888
25				16, 645	16. 900	8. 560	17.840	8.090	20.149
	PLACE 1011407	20.446	18. 260			4. 330	7.245	6. 219	8.740
	PLACE 1011419	9.047	8.378	6, 933	5. 544			15, 534	27. 379
	PLACE 1011433	13, 904	35, 637	21.499	14. 088	12.024	17. 492		
	PLACE 1011440	57.799	31.667	21.664	18. 327	13.093	34. 588	30.019	22, 159
	PLACE 1011452	50.007	42.314	37, 053	49. 949	18.696	31.802	28.114	34.843
	PLACE 1011465	35. 426	19.398	13.047	12.250	12. 486	25.628	23. 462	18.107
30	PLACE 1011472	62.882	51, 139	24. 865	13.679	29. 181	24.440	22. 986	20.138
	PLACE 1011477	56.690	73.733	72. 345	49.100	38. 345	43.680	52.566	88.177
	PLACE1011478	63.612	53.418	38. 381	43.231	28. 020	32.283	28. 922	47.558
	PLACE1011492	106.290	57.337	44.835	33.949	26. 366	41.775	46.645	28.355
	PLACE 1011498	11.479	10.039	1.690	3.014	1.593	3.917	8.921	0.000
	PLACE 1011501	6.078	13. 915	3. 925	4. 468	3. 927	10.819	15.717	55.041
35					0.243	2.449	3.045	3.606	2.018
00	PLACE 1011503	1.874	0.762	1.380			9.010	13, 173	
	PLACE1011509	15, 310	13.049	7.406	5, 231	8. 198			13.881
	PLACE1011514	63.158	72.840	43.610	53. 595	30.828	44.567	49. 208	51.604
	PLACE1011518	26.859	55.632	40, 993	27.965	33. 580	27.829	35. 366	35. 955
	PLACE 101 1520	4,008	12.681	4, 680	4.882	2.815	4.425	5.052	6.373
	PLACE 1011538	46. 942	112.381	14. 535	10.906	7. 023	21.261	18. 123	15, 061
40	PLACE 1011555	64.949	. 24. 945	16,779	8.387	10.043	27.860	31.802	9. 584
	PLACE 1011561	10.363	15. 824	6.531	16.410	4.737	8.801	9. 321	17.672
	PLACE 1011563	10, 025	6, 203	5. 528	4.965	4. 378	7.900	10.397	8.513
	PLACE 1011567	42. 901	33, 701	15. 168	22, 187	13.471	15.650	16.469	24.618
	PLACE 1011569	26. 547	51.848	39.883	37. 100	23. 589	23. 252	34. 227	41.438
	PLACE 101 1578		90. 143	56,009	77.009	47. 187	46.612	36. 385	75. 351
		65. 455				15. 477	22.594	28.411	25.540
45	PLACE 1011586	46.138	39. 212	16,045	20. 957				
	PLACE 1011635	16.794	16.170	6.079	7.918	5. 168	11.027	22.021	10.224
	PLACE 1011641	1.228	0.000	3, 690	2.905	1, 954	3.104	3.300	2. 256
	PLACE 1011642	17.749	23. 124	9. 273	20. 132	5.674	11.138	15. 685	20.899
	PLACE 1011643	26. 441	17. 121	11.726	11.897	5. 398	10.061	15. 157	16. 472
	PLACE 1011646	84. 129	76.809	63. 483	58. 487	61.819	45.212	52.514	53.689
50	PLACE 1011649	148, 652	79.404	41.401	24.880	37.816	60.892	98.048	59. 957
50	PLACE 1011650	207.033	106, 793	62, 104	33.902	59,773	85.346	101.285	59.651
	PLACE 1011661	89. 284	69, 963	52.044	60, 130		46.476	38.780	47.335
	PLACE 1011564	19.831	24. 910	9.719	12.162	10. 285	14. 197	16.087	9.849
	PLACE 1011672		4. 324	0.000		2.518	4.317	5. 108	6.001
		3. 166					13.639	14. 193	4. 640
	PLACE 1011575	5. 381	4. 183	13.639	3.525	18.043	1 13.033	1 14. 133	7.040

561

Table 152

				Tab	le 152				
	PLACE1011682	46.195	19.920	15.150	18. 241	16.697	20.650	33, 169	20.683
	PLACE1011708	140.868	80.025	46.997	44. 349	48.806	85.376	98.779	53.876
	PLACE1011719	81.308	62.978	42.651	25. 199	32.975	36.215	54. 409	40.754
5	PLACE 101 1725	51.825	51, 140	27.931	38.736	21.984	25.006	26. 264	40, 599
	PLACE 1011729	24. 560	24.476	13.172	17.322	11. 225	10.549	10. 437	11.441
	PLACE 1011741	10,084	12.651	9. 857	10.562	8.885	9.463	12.550	13.970
	PLACE1011749	65. 367	64.514	37. 914	40.516	34.378	23.889	21.546	32.149
	PLACE1011757	18.814	44, 445	37.496	28. 407	37.470	13.419	20. 349	44. 087
	PLACE1011762	22, 509	23.571	12.319	14.785	13.545	12.246	16.007	14.719
10	PLACE1011778	18.861	10.736	11.124	6.662	7.815	9.039	11.917	6.723
	PLACE1011783	121.850	129.976	50. 595	57. 237	55. 572	43.090	130, 253	61.954
	PLACE 101 1795	31.927	47.460	15.112	14. 530	14.324	16.899	13. 987	13.824
	PLACE 101 1810	11.913	20.873	9.752	5. 145	8.850	7.953	21.005	8. 397
	PLACE1011824	19,075	38. 642	12.337	13. 272	9, 167	11.037	20.832	9.083
	PLACE1011825	101.516	76.411	46.000	26.850	59, 669	37.495	57.769	32.550
15	PLACE1011835	41.770	35.699	13.510	12.484	12.451	13.661	25. 449	15. 527
	PLACE1011836	75.164	61.584	46.814	31.866	60.375	30.344	47.168	42.711
	PLACE1011847	13.876	13.405	4. 281	8.038	4. 394	3.642	10.641	8.968
	PLACE 101 1855	23, 160	24.900,	11.611	9. 421	10.774	9. 353	18. 255	8. 246
	PLACE1011858	17.703	19.170	8. 339	6. 242	7. 166	9. 321	11.444	10.044
	PLACE 1011874	25. 436	29.797	26.222	32. 382	13.428	18.138	15.516 8.980	21.195 5.065
20	PLACE 1011875	3.069	12.743	6.998	4. 382	6. 338	8. 026 26. 880	25.751	19.376
	PLACE1011877 -	32.981	22.725	17.384	15.505	5.675		35. 929	22, 229
	PLACE 101 1891	49.673	22.359	23.890	9.835	15.099	27.985 0.000	6.891	3.826
	PLACE1011896	4. 107	0.000	3.756	17, 707	11.558	18.630	38. 456	21.819
	PLACE 10 1 1920	31.343	26.346	21.681	29.603	0.000	23.870	31.601	34, 831
	PLACE 1011922	42.691 32.608	40.664	19.701	8.083	15. 625	16,742	22.157	29.615
25	PLACE1011923 PLACE1011937	92.606	35.417	26. 508	20.596	9. 785	43.673	39. 451	0.000
	PLACE1011939	84. 529	52.763	38. 555	19.964	7. 336	39.880	62.062	31.494
	PLACE 1011940	59.607	59.623	43, 124	27.246	23.603	35. 438	69.861	53. 973
	PLACE1011962	100.298	63.747	55.070	41.766	37.368	61.832	65.091	63. 896
	PLACE 1011964	11.886	16.598	13.946	17. 132	. 12. 848	11, 456	26. 353	18. 276.
	PLACE1011978	18.640	19.836	21.517	38. 291	0.000	21.287	15.757	50, 491
30	PLACE1011980	92.462	82.334	53. 193	72.449	39, 473	41.547	40.407	54. 365
	PLACE 1011981	61.362	58.174	46.817	28.272	28. 476	43. 347	64.658	50.398
	PLACE 1011982	15.790	14.181	4.817	8.312	3, 604	7.757	7. 260	0.000
	PLACE 1011995	86.516	35.794	56.068	64.038	31.871	35. 426	30.449	14.091
	PLACE1012023	13.104	15.527	8.953	7.883	5. 966	11.716	8. 264	5.743
	PLACE 1012026	7.250	6.837	6. 369	2.909	2.441	4.999 7.545	11.516	13.665
35	PLACE 1012031	17. 346	7.096	7.365	6, 293	92.061	81.725	104. 515	91. 349
	PLACE2000003	208. 422	130.772	108.228	19.141	15. 235	28.560	47. 298	41, 315
	PLACE2000005	71.165 39.195	33.762	22.805		19. 193	16. 829	26.310	26. 260
	PLACE2000006 PLACE2000007	49. 369	22.909	15.022	10.283	10.043	26, 310	24.168	17.472
	PLACE2000011	71. 136	45. 914	39.612	33.759	26.056	33.405	30.793	16.938
40	PLACE2000014	10.718	21, 905	13.060		8. 179	11.383	26.861	29.837
40	PLACE2000015	5. 458	4.184	2. 923		2.593	2.078	3. 383	5. 945
	PLACE2000017	46. 332	45.480	23.941		21.386	18.932	16.284	17. 911
	PLACE2000021	17, 344	18.232	12, 294	30.435	15. 289	14.854	16, 815	25. 461
	PLACE2000022	214.445	144.482	60.979	80.113	67.083	66.854	70.170	73.024
	PLACE2000030	187.619	114.314	63, 549			68. 183	115.701	63, 549
45	PLACE2000032	87.441	77.188				33.214	31.270	38. 239
43	PLACE2000033	19.139	24.471	9. 846				9. 886	
	PLACE2000034	42.847						25. 238	
	PLACE2000039	132. 992					60.606	56.917	66. 559
	PLACE2000043	79.648					29.880	42.418	
	PLACE2000044	108. 910						73.844	
50	PLACE2000047	152.880	109.530						
50	PLACE2000050	152, 213							
	PLACE2000061	29.004							
	PLACE2000062	72.911							
	PLACE2000072	26, 412							
	PLACE2000073	30, 538	11.955	9, 19	7   2.761	2.738	11.625	16.675	1.333

562

Table 153

						77.77	20 000	24 210 1	26.522
	PLACE2000097	26.855	20.822	13.598	19.129	11.744	36.567	24.316	
	PLACE2000100	65.222	58.680	32,787	36.772	34.459	33.979	29.850	45. 558
		87.537	67.579	43.315	48.791	32.811	36.573	42.226	40.018
	PLACE2000103								50.310
5	PLACE2000106	109.631	86.434	50.857	64. 247	33.823	50.406	61.798	
	PLACE2000111	67.743	54.614	49.948	32.461	25.661	39. 498	39.424	45. 128
				16, 909	7.307	8.764	18. 404	25.904	9.246
	PLACE2000115	39.616	21.252						198.972
	PLACE 2000118	525. 051	269.098	228.675	184.616	169. 233	347. 407	255.751	
	PLACE2000124	349. 581	275.812	246.822	261.885	198. 107	204.514	181.925	195.861
						44. 528	98.585	135.305	44.096
	PLACE2000132	219.428	75.779	55, 477	27.845				16.417
10	PLACE2000136	26.471	13.700	10. 948	5. 229	9. 552	12. 386	20. 372	
	PLACE2000137	136. 105	55. 207	38.965	25. 263	38.081	60. 138	77.269	40. 309
			58. 228	35. 563	26.913	23.042	43. 520	44. 482	46.587
	PLACE2000140	61.894					15. 456	20.704	13.852
	PLACE2000147	35.744	28.047	17. 366	9. 287	7.856			
	PLACE2000153	10.251	5.138	4.944	3. 289	1. 583	8.639	8. 187	4.258
		28. 952	20.099	15, 192	12.672	6. 324	15. 972	21.497	18.781
	PLACE2000164					22.805	27.949	31.835	29.350
15	PLACE2000170	59.457	56.458	26.480	33. 136				
	PLACE2000172	44. 931	19, 156	12.587	7. 529	12.190	15. 161	26.980	18.906
	PLACE2000173	61.374	67.180	25, 374	32.768	28. 635	32. 143	41.210	43.509
						30.132	26.716	42.849	35.942
	PLACE2000174	58.350	40.462,	27.593	30.601				
	PLACE2000176	67.823	54.888	28.038	22. 906	22.587	33.838	40.703	24.007
	PLACE2000187	58. 492	46.505	35,000	29.053	17.412	35. 409	39.960	33.655
				34. 386	16.556	25.028	35.589	41.068	25.755
20	PLACE2000216	67.045	48.042				45. 597	40.342	27. 793
	PLACE2000219	102.450	53.525	43.919	40.723	27. 590			
	PLACE2000221	172.504	104.236	71.274	95.080	69.068	73.974	75.780	84. 353
	PLACE2000223	1. 924	0.000	0.337	0.072	0.489	0.000	1.615	0.884
					6.372	. 14, 358	19.848	38.853	25. 391
	PLACE2000231	46.085	20.513	17.117					
	PLACE2000235	124. 328	101.132	67. 369	85.561	58. 141	54. 197	68.871	76.828
	PLACE2000246	104.336	91.568	43. 204	38.961	37.372	49. 589	53.726	38.556
25		80, 119	58.341	26. 725	32.576	29. 924	31.634	35, 536	46.483
	PLACE2000264					44.169	82.504	117.862	37, 316
	PLACE2000274	178.113	50.862	46.488	15.876				
	PLACE2000287	132.856	101.370	60.760	60.336	35. 602	65.837	81.297	70.741
	PLACE2000296	49. 120	36.473	16, 163	15.750	18. 250	18. 313	35.709	33.015
					27.707	22. 845	20.720	30, 271	32.036
	PLACE2000302	57. 145	42.035	23. 159					
	PLACE2000305	175.494	200.830	97.799	110.854	103. 121	82.645	117.383	111.334
30	PLACE2000317	43.989	47.859	17,789	17.969	19.049	22.044	50,064	40.575
		0.000	7.097	5.063	2.422	6. 266	5.462	10.248	7.127
	PLACE2000324					19. 395	41.882	61.667	32.402
	PLACE2000334	68. 183	58. 423	27.660	13.890				
	PLACE2000335	124.754	148.141	79.507	92.542	69. 951	70.762	73.634	60.750
	PLACE2000340	26.477	26.590	14.223	111.260	9. 640	12.040	23.150	14.154
					25.403	26,509	38.745	65. 207	46.925
	PLACE2000341	77.833	55.873	31.663					
35	PLACE2000342	106.364	52.711	44.616	37.588	44. 103	48.901	80.862	45.540
	PLACE2000347	135. 574	132,050	56, 804	42. 203	56. 182	70.882	92.167	64.861
		93.053	95.338	40.039	30.886	41.634	43.514	108.320	65.738
	PLACE2000357						30.828	46.656	38.072
	PLACE2000358	37.940	54.020	19, 892	24.091	19.855			
	PLACE2000359	44.601	31. 382	22. 450	28.212	15.793	15, 150	23.074	23. 575
	PLACE2000366	121. 162	103.772	44,748	43.347	42. 993	37.451	42. 382	49.575
	PLACE2000371	30. 423	16.028	14.211	9.577	16.570	13.288	16.943	9.168
40						27. 244	41.111	69,708	39.196
	PLACE2000373	103. 200	59. 241	36.611	28.313				
	PLACE2000374	113.892	55. 366	30. 642	21.105	30, 506	39.759	73, 431	35.604
	PLACE2000379	20.349	15. 495	7.621	6.080	7. 432	5.799	10.929	11.275
	PLACE2000386	2957. 979	598. 564	744. 423	192.993	914. 385	1779.750	2073. 338	474.610
							36.369	40.361	36. 550
	PLACE2000388	71.861	48. 309	26.919	20, 159	20.978			
	PLACE2000392	352. 525	278, 976	168.585	149. 394	126.536	186.631	228, 238	160.402
45	PLACE2000394	53.696		49.507	50. 392	15. 244	41. 226	40, 124	40, 112
	PLACE2000398	108.135		58.643					
	PLACE2000399	67.901	42.851	38, 688	28.243		41. 332	58, 492	
	PLACE2000402	63. 927					39, 649	49, 188	31.169
							36, 900	57.711	32.512
	PLACE2000404	52.116							
	PLACE2000411	344.233	265, 387	148.539	150.545	127.069	193.357	280.999	
50	PLACE2000418	98. 999					47, 480	51, 418	37.707
							86.077	101.959	
	PLACE2000419	173.685							
	PLACE2000425	48.498	43.030	24.787	27.067	15.782	34.775	41.783	
	PLACE2000427						40.769	48.213	24.735
	PLACE2000433	85.693	46.037	39.587	31.830	1. 65. 130	1 41. 363	1 43, 179	1 99.010

563

Table 154

							262 644	106.234
PLACE2000435	627.805	144. 999		57. 999	150.754			
PLACE2000438	56.718	23.072	24.569	11.907	16.555	28. 249	47.594	27.160
PLACE 2000450	154.687	141.268	71.263 1	18.445	53.787	64.455	51.630	74.221
PLACE 2000455	67.470	36. 100	20.827	16. 588	21.358	35.782	37.471	28.970
PLACE 2000/458	104. 672	42.860	43.528	21. 379	21.800	58.644	70. 147	47.611
PLACE2000464	105. 901	34. 595	41, 144	17. 313	27, 129	60.407	69.472	24. 244
PLACE2000465	80, 401	104. 292	71.810	73.398	40, 246	50.872	47.384	76. 283
	420.021			62,099	161.932	255. 494	273.349	340. 926
PLACE2000473			4. 305	5. 801	6.451	4.611	5. 267	7, 301
PLACE2000477	15.988	4.741	55.053	66.698	49.843	55. 946	72.642	62.012
PLACE 3000004	150. 291	87.960		37.661		010.865	903. 209	416.227
	2308.534				45, 276	64.510	77. 399	65. 319
PLACE 3000020	129, 151	97.914	62.838	38.547	20, 999	29. 355	27.242	35.815
PLACE3000029	65.003	63.572	28. 269	34.591		26.851	33, 481	34. 526
PLACE 3000038	60.832	39. 883	31.082	31.184	18.004		44. 374	46, 348
PLACE 3000052	80.986	57. 505	42.010	29.980	28.396	38.406		
PLACE 3000059	14.309	10.723	7.978	9.607	8.219	10.311	10. 168	14. 892
PLACE3000067	148.633	122.359	72.812	107.464	67.921	73.383	51. 333	59.832
PLACE3000069	94.472	58, 891	39. 254	38.134	38.532	45.365	50.569	49.095
PLACE 3000070	606.923	398.146,	277.096	302.922	122.053	340. 430	303. 517	264, 024
PLACE 3000103	37.665	49. 384	23.681	21.788	14.319	19.018	24, 128	23.022
PLACE3000103	71.233	77.814	32.703	36.829	29.100	33, 236	32. 954	38. 558
PLACE3000121	28.770	15.821	15.686	6.769	11.774	14.597	21.083	15.791
PLACE 3000124	136.225	102.926	73.102	88.816	50.098	64.249	70. 972	88.890
PLACE 3000135	5. 325	1.538	1.703	2.303	1.771	1.846	2.243	1.568
PLACE 3000 135	264.467	146.748	117.350	79.000	94.116	149, 983	135. 199	82.816
PLACE 3000 142	84. 493	43.724	33.445	21.753	29.470	39.958	59.408	36.420
PLACE 3000142	202.991	105. 472	78. 043	43. 347	67.611	96.794	127.254	104.646
	45. 022	53.334	28. 294	29.723	15.237	24, 991	20. 260	28,896
PLACE 3000147	50. 238	25. 306	12.752	14, 405	11.331	16.047	21.617	14. 375
PLACE 3000 148			5. 983	4.462	5, 238	17.888	19, 855	5.659
PLACE3000154	16.588	24. 537		61.895	55.036	69,498	99.138	67.036
PLACE 3000155	162.823	103. 374	73. 169	36.695	86.574	251.934	180, 898	69.146
PLACE 3000 156	293.645	80.486	96. 151	23.067	21,480	31, 175	45. 472	36,779
PLACE 3000 157	77.274	48. 353	30. 271			58. 354	55. 566	83. 250
PLACE3000158	138.262	117.084	66.013	76.854	56.610	7.038	7.949	11, 165
PLACE3000160	12.383	13.802	3. 360	3.545	2.772	41.506	37.316	64. 578
PLACE3000169	112.273	107.072	60.628	74.727	35.758	112,925	84, 309	50, 931
PLACE3000181	159, 980	52.030	56.098	26.437	39.138			36.285
PLACE3000194	59. 243	40.406	43.072	30. 599	27.793	33.533	39, 940 96, 254	57.504
PLACE3000197	2.773	2.051	1.429	3.753	0.000	3,916		12.260
PLACE3000199	38. 435	22.543	11.795	7.257	11.967	16.257	14.819	65.731
PLACE3000205	98.788	82.371	76.207	41.507	69.168	50, 577	62.634	
PLACE3000207	107.828	91, 992	61.336	61.872	58. 924	42.359	53. 327	75. 106
PLACE3000208	. 112.570	54.203	55. 951	38.351	49.935	44, 990	75. 532	53. 240
PLACE3000213	26.219	39.836	11,741	11.345	7.948	12.842	24. 022	17.439
PLACE3000215	90.876	34.688	28, 635	9.043	15. 498	40.462	43.681	18.877
PLACE3000218	10.221	2.943	2.894	3.797	1.404	4.853	5.114	3.490
PLACE3000220	61.519	52.284	29.152	23, 405	20.917	20.102	32.078	28. 959
PLACE3000221	57, 492	57.641	28.073	44, 309		41.840	33, 858	52. 488
PLACE3000225	73.279	54. 393	35. 962	36.879	33, 401	26.367	40.176	43.907
PLACE 3000226	73.816	45.891	30. 595	22.786	30.642	32.460	45, 062	32. 422
PLACE 3000230	46.785	26. 306	16.545	6.639		18.992	43.959	26.308
PLACE3000231	48. 528	32.588	17. 433	13.571	12.141	20.113		18, 127
PLACE3000235	85.027	89.322	36, 118	40.285	33.985	29.150	33.828	45. 276
PLACE3000242	40. 499	25.236	19, 477			22.181	24.892	16. 933
PLACE3000244	8.374	6. 431	4, 114		1.774	5.910	8. 022	3.080
PLACE3000253	15, 620		14,659			14.844		13.77
PLACE3000254	1079, 768	504. 372	199.997					328. 912
PLACE3000234	142.610	130.398	184, 934					90.942
PLACE3000276	50. 360		20. 928					23.25
	753.417		316, 576					311.942
PLACE3000304			22.694					66. 550
PLACE3000309			5. 349					
PLACE3000310			10, 590					
PLACE3000320			19. 280					
PLACE3000322	59. 027	28.943	13.200	21.430	1 23.010			

Table 155

	101 105 1000110	216 200	112 224 1	67.450	57. 211	77, 413	103.940	153.059	92.002
	PLACE 3000 330	216.369	112.324						
	PLACE 3000331	175. 154	109.366	65. 174	61.976	64. 453	62.935	100. 525	16.877
	PLACE 3000336	72.694	51.382	24.596	19. 447	23.969	28.997	69.879	41.462
5	PLACE 3000339	30,681	21, 404	10.699	10, 293	10.726	16.029	27. 155	16.949
			47.003	23.728	24. 494	19.963	23.504	27.314	25.967
	PLACE3000341	60.229						11. 370	19. 246
	PLACE3000350	29.438	30.806	13.412	18.894	10.478	10.194		
	PLACE 3000352	133.033	66.842	27. 124	22. 155	31.563	39.432	49.060	26.130
	PLACE 3000353	43.758	27.987	13.231	12.753	11. 266	21.242	42.590	23.677
	PLACE 3000 362	67.720	68.906	37.576	54. 837	39.546	31.556	42, 322	53, 834
10					14. 578	19.607	23.664	45.846	21.687
10	PLACE 3000363	57.403	38.780	25. 263				58. 355	39.738
	PLACE 3000365	67.367	70.204	26. 265	27. 858	28.782	28.393		
	PLACE 3000373	13.237	14.898	8. 160	10.479	10.212	7.337	11.629	5. 545
	PLACE 3000374	65.194	47.989	28. 255	34. 215	25.888	25.506	40.045	29.881
	PLACE3000387	39.123	14.751	9. 548	6. 520	10.023	13.134	24. 323	11.700
					30.962	16.063	16.391	26.317	35. 550.
	PLACE3000388	38.498	49.657	25.044					
15	PLACE3000399	148. 163	127. 490	65. 532	74. 992	56.760	50.436	71.879	57. 275
	PLACE3000400	64.113	49.775	24.696	24. 323	28.318	34.732	29. 297	29.946
	PLACE 3000401	643.361	789.055	443.841	553. 459	428.754	314.650	347. 522	356.250
	PLACE3000402	93. 152	75.383,	36.033	35, 535	33.800	26.510	39, 162	39.094
						23.948	53.017	69.999	41. 988
	PLACE 3000405	116.575	74.775	47. 203	35. 397				
	PLACE 3000406	46.734	47.216	28. 404	38. 943	18.564	21.735	18. 510	22. 439
20	PLACE 3000413	172.089	63.768	60.797	25. 154	38.861	86.736	100. 967	39. 294
20	PLACE3000416	72.812	94.541	27.443	24. 126	23.401	36.001	52.778	31.746
	PLACE3000425	75. 299	85. 243	55.831	51.775	29.832	40.724	54, 413	42.424
						53.901	88.235	140.605	71.376
	PLACE3000437	152.596	106.131	91.713	79.520				
	PLACE 3000455	199.980	144.915	86.941	70.024	46.162	89.704	140.865	87. 299
	PLACE 3000475	344.660	151.608	142.664	51.432	168. 147	291.157	322.276	96.413
	PLACE3000477	105. 902	72.097	35.966	23.877	17.322	48.569	53.837	34.942
25	PLACE 4000003	21.542	6.768	7.756	4. 338	6.666	15, 322	9,008	10.517
					29. 174	47.030	40.800	60.131	38, 137
	PLACE4000008	81.624	76.594	49. 347			123.118	127. 853	
	PLACE 4000009	254. 207	142.614	81.374	67.480	67.588			80.493
	PLACE4000014	93.227	49.366	32.322	19.702	25.748	40.531	72.266	36.731
	PLACE 4000029	21.650	25.863	17.118	20.048	17.456	36.449	43.230	27.051
	PLACE 4000034	49, 161	79.725	28. 534	23. 533	19.403	43.040	40.269	23.537
30		166.916		69.807	85. 324	50.891	74, 119	64, 317	64, 497
••	PLACE4000049		134, 169				36, 433	50. 349	23. 477
	PLACE4000052	54.863	57.074	25. 752	30. 034	15.812			
	PLACE4000062	78.176	55.581	32.501	23. 565	14.025	47.511	74.636	26.040
	PLACE 4000063	84.945	48.380	39.855	15.974	28.354	50.659	60. 330	32. 588
	PLACE4000089	19.057	35.752	29, 230	17.534	17.492	11.406	15.833	13, 554
	PLACE4000093	26.060	15.272	12.061	6.706	13.618	11.634	18.344	13,777
05							33. 159	29.333	44.906
35	PLACE4000100	101.893	42.734	31.255	36, 161	14.062			
	PLACE4000103	124, 173	34.660	22.754	19.690	20.649	30.763	70.971	19.503
	PLACE 4000106	98.597	75.194	36.209	29.412	33.084	61.638	76.538	44.570
	PLACE4000128	129.329	131.483	60.440	57.978	41. 117	68.736	84. 185	95. 597
	PLACE4000129	132.932	37.431	53. 267	53.097	33.745	72.527	81.857	45.648
	PLACE 4000131	156. 165	156. 169	86.886	106.633	78.888	107.180	102. 299	56.814
							5. 146	10.290	4.656
40	PLACE4000147	16.492	9.413	7. 966	2. 107	5.770			
	PLACE 4000156	69.314	72.955	65.884	87.221	44.343	45.822	35.362	77.048
	PLACE4000175	60.994	54.028	16.876	13.509	17.492	17.584	32.845	26.309
	PLACE 4000190	593.634	220.190	171.592	116.664	189.541	260.140	310.147	138.653
	PLACE 4000192	301.266	121.069	80.280	70.432	67.302	127.637	134. 475	72.627
	PLACE 4000208	259.054	236.436	119.680	97.518	77.872	86.994	97.682	154, 492
							149.275	122.028	95. 561
45	PLACE4000211	242. 387	150,657	98,746	66.861	74.283			
	PLACE4000214	67.058	61.229	37.510	23.741	22.459	30.120	39.510	34, 868
	PLACE4000222	106.945	86.369	43.808	41.733	40.284			46.046
	PLACE4000223	107.887					37.870	44.089	22. 256
	PLACE 4000229	50.488					27.323	36.955	19.875
							28.713	41.794	26. 235
	PLACE4000230	83.847							
50	PLACE4000233	96.059		60.661				40.716	47.823
50	PLACE4000239	124. 398	94. 107	57.093	48.109	34. 394	43.667	35.791	41.364
	PLACE4000247	54.958					25. 144	27. 546	21.593
	PLACE4000250	104, 404						60. 153	65.709
								16.958	13.472
	PLACE4000252								
	PLACE4000259	113.573	49.555	27.075	21.856	43.353	30.044	52.944	26.431

Table 156.

					16 130.				32 888 T
	PLACE4000261	254.068	48.744	84.359	22.460			113.583	36.809
	PLACE4000264	39, 731	29.931	13, 801	9. 433	14. 239	13.997	27.405	15.510
	DI 165 4000504	85, 391	69, 167	59, 645	32.049	32.560	45.023	59.556	44.048
	PLACE4000269		35. 516	21.356	20.188	22.145	14.735	20. 334	24,680
5	PLACE4000270	37. 293			24.956	58. 105	75. 320	65, 805	106.301
	PLACE4090281	*****	130.790	50, 704	44.034	42.272	39,616	55.059	49.678
	PLACE4000300	95.228	64, 001			37. 273	44. 289	54. 376	57.927
	PLACE4000320	101, 920	74.756	53.518	50.074			67. 236	63.467
	PLACE4000323	106.246	90.568	59. 225	75.643	65. 195	71. 824		24. 334
	PLACE4000326	50.786	39.408	21. 110	15.693	16. 385	24. 171	25.892	
	PLACE4000344	47. 237	25.071	19.282	9.754	15. 816	17.064	28. 344	22.605
10		270.519	135. 102	97.629	73.164	79.089	145.628	174.326	132.718
	PLACE4000347		69.949	21. 125	14, 137	10. 506	24.887	36.668	32.881
	PLACE4000354	51.402		13. 300	12.406	12. 328	16. 184	16.983	12.631
	PLACE 4000367	38. 537	21.917		18, 841	17. 942	39.036	46.668	28.559
	PLACE4000369	87.562	48.818	27. 044		28. 043	28, 363	34. 411	32,783
	PLACE4000379	63.427	45.050	34. 549	40.613			27.432	22.848
	PLACE 4000387	51.546	28.804	20, 204	18.439	20. 155	20. 584		
15	PLACE 4000392	16, 062	7.012	6, 606	5. 828	5.717	7. 153	9.447	4.556
	PLACE4000399	537. 973	347.563	215.840	188. 160	226.834	294, 013	378.985	282.393
	PLACE 4000401	18.633	16.086	12.450	8.891	4.760	9. 336	9. 594	9.016
		122.680	74.783	64. 480	32.311	31.018	61.677	74.741	57.710
	PLACE4000403		69.288	26.062	27. 151	18.908	24.969	28.090	28.450
	PLACE 4000411	76.474		42.067	13.307	25. 782	58.009	67. 901	23.594
20	PLACE4000415	117.128	42.809		24. 312	34. 852	60. 268	78. 927	60.597
20	PLACE4000416	155. 173	151.945	41.224		15. 430	26. 353	51, 392	21.253
	PLACE4000424	49.737	20.818	19, 113	10.882		23. 575	63.847	39.828
	PLACE4000431	94, 197	46.298	22.172	18.259	30.613			4, 152
	PLACE 4000443	5.628	10.390	1.885	3.662	4. 723	4. 338	8.728	
	PLACE4000445	112.063	123.064	82.212	73.969	75.667	71.847	80.872	98. 196
	PLACE 4000450	235.301	129.164	80.479	\$8.100	59.886	126. 244	134.749	85.784
25	PLACE4000455	48, 423	52,624	22.324	12.728	17.652	29. 121	33.876	28.299
23	OLACE 4000435	106.018	96.543	76.272	77.100	59.155	46.270	60,646	57.534
	PLACE 4000465		313.894	141.390	142.098	110.817	145, 538	179.778	235. 989
	PLACE4000466	291.255		184.390	172.988	162.349	205. 973	249.573	175.977
	PLACE4000472	361.477	283.612		34, 491	38. 357	27.786	47.292	39, 254
	PLACE4000487	71.130	60.554	31.674		29, 344	31.561	68. 977	55.815
	PLACE4000489	95, 437	42.543	25.117	24. 559	26.845	35.819	41.938	46, 527
30	PLACE4000494	88.573	62.176	35, 502	19.031			78. 120	106.317
	PLACE4000502	149.633	181.173	61.673	64. 434	54. 907	64.869		41.190
	PLACE4000521	204.368	58.842	53.769	22.018	19. 396	90.039	90. 251	
	PLACE4000522	70.773	56.092	27.371	16.069	23.518	31.461	43. 466	39.760
	PLACE4000537	155. 193	45.421	44, 392	17.892	44. 281	65, 488	98.332	46.179
	PLACE 4000548	47.086	28. 598	16,763	16.406	16.740	16.619	38.465	30.778
		10.369	12.539	7.971	5.855	5. 400	4.652	8. 570	10.740
35 ·	PLACE4000558		51. 427	22.039	21.955	29.024	23.682	53.726	32.562
	PLACE4000581	70.383		5.754	7.501	7, 952	10.260	10.943	10.189
	PLACE4000590	24.623	8.914			26.365	31.598	47.539	27, 961
	PLACE4000593	72.087	47.632	23.074	21.723		178. 284	193.620	70.237
	PLACE 4000612	363.116	155.910	113. BOO		124. 093	34.764	51. 100	28. 946
	PLACE 4000638	77.534	58. 517	30.744		38. 112		43.714	24. 670
	PLACE4000650	45. 331	35.490			17.671	20.345		
40	PLACE 4000651	81.785	55.336			31, 108	38.514	81.922	45. 304
	PLACE4000654	6. 383	10.852	2.069	2.695		0.000	8.009	5.077
	PLACE 4000670	26.614			5.853	8. 977	8.517	8.611	9, 175
	PLACE4000685	353, 509					251.552	212.919	154.500
	PLACEHOUGES	6.072					6.156	75.595	9.677
	PLACE4000687						21.575	38.678	19.632
	PLACESO00003	40.413					14,637	23.435	12.833
45	PLACE5000005	29. 397	16.490						
	PLACE5000019	23. 138							
	PLACE5000021	11.535	7.579	5. 66					
	PLACE5000022	46.567	29.719	16.48	17,005				
	PLACE5000024					17. 296	33. 257		
								49.655	22.062
	PLACE5000036								
50	PLACE 5000059								
	PLACE5000076								
	PLACE 5000117	42.649							
	PLACE5000143		38. 12	4 31.38	8 29.11				
	PLACE5000152			3 4.88	0 1.27	8 1.829	5.71	9. 92	4, 547
	F LACE 3000 132		- 1 3 - 1						

Table 157

	Gi verrene	70 004 1	25 002 1	31 836 1	AC CAE 1	17 671 (	22 026	42 636 1	45 054
	PLACE5000154	70.894	26. 982	21.228	26.625	17.971	32. 836	43.570	45.054
	PLACE5000155	443.969	270.563		139, 163	137.024	244. 271	195.771	169.360
_	PLACE5000165	529. 207	254.686		123.963	145. 432	257. 836	242.614	165.245
5	SKNMC1000004	20.836	13. 305	17.789	33. 557	11.594	10.964	6.648	21. 308
	SKNMC1000011	19.687	9.046	-7. 372	8. 263	7.296	15.689	11. 182	14.777
	SKNMC1000013	9.401	12.821	9. 287	3.794	5.931	6. 702	10. 997	8.736
	SKNMC1000014	49.003	43.832	32,008	24. 581	23.480	20.065	21. 197	18,671
	SKNMC1000018	33.522	17.298	13.017	4. 235	8.795	16. 555	20.822	15.790
	SKNMC1000020	41.784	25. 172	10.947	6.057	6.258	17.499	22. 243	15. 547
10	SKNMC1000045	21.429	19.675	15.389	7.367	8.974	13. 224	14, 566	12.097
	SKNMC1000050	22.145	28.518	10.065	7.977	7.275	14.859	11.644	8.042
	SKNWC1000062	338. 427	274.434	175.123	132.052	150.251	235. 537	155.269	137, 370
	SKNMC1000075	20.756	21.072	10.730	10.756	8.063	10.684	15. 925	10.454
	SKNWC1000082	24.604	10.450	9.435	7.978	7.660	10,818	12.376	12.685
	SKNMC1000091	36.258	20. 984	12.691	12.987	9.671	18, 161	17.028	15.150
	SKHWC1000091	27.554	15.672	10.331	8. 117	8.086	17.003	23.741	6. 484
15				9.892	7. 092	9. 487	18.879	22.259	6.010
	SKNUC1000104	38.010	34. 379			15.067	12.794	17.603	10. 906
	SKNMC1000113	39.920	26.152	14, 548	11.762		35. 214	34.073	39.116
	SKNMC1000119	68.128	70. 122	43.005	35. 267	28.955			
	SKNMC1000142	32.190	14.734	11.314	9.644	8.615	13.750	11. 275	11.126
	SKNMC1000170	27.877	27.618	13.752	9.407	7.172	15, 123	19.813	13.284
20	SKNMC1000178	70.066	63.234	33,059	29.079	25.498	40. 509	40.085	31.660
	SKNMC1000194	49.613	30.075	14. 523	13.545	13, 410	19. 965	25. 730	19.940
	SKNMC1000198	36.190	10.269	18. 321	16.365	19.849	22. 261	23.973	21.923
	SKNWC1000225	20.577	23.995	7.702	12.589	11.016	9. 595	24.700	19.639
	SKNMC1000249	35.318	7. 307	2.999	2.393	1.501	10.815	6.991	7.735
	SPLEN1000007	17. 285	35.392	16.709	18.674	6.880	10.787	9.808	21.699
	SPLEN1000012	79.902	26.456	22.780	18.019	22.231	32.118	34.361	44. 355
25	SPLEN1000014	86.560	12.587	39.565	11.907	15.132	29.061	14, 109	26, 990
	SPLEN1000036	39.586	28.908	15.910	11. 331	10.780	20. 946	21.977	20.383
	SPLEN1000059	0.000	0.000	0.000	0.000	0.000	0.000	0.000	2.428
	SPLEN1000068	42.216	61.333	16.982	24, 191	14.315	17.337	19.640	62.286
	SPLEN1000072	80.933	51, 171	18.946	13. 264	24.375	35. 212	43.609	27.418
	SPLEN1000101	56.109	102.035	38.061	51. 936	36.704	44. 974	32.451	43.746
30	SPLEN1000108	28.462	16.640	8.555	5.187	16.134	11, 421	13.414	8.735
	SPLEN1000113	51.510	25.822	25.943	12.537	11.070	26.855	29.899	18.889
	SPLEN1000114	35.034	24. 235	14.342	6.652	10.171	15.802	22.089	21.376
	SPLEN1000132	49.855	38.464	20.708	19.052	11.964	26.849	30.806	43.790
	SPLEN1000135	69.620	36.735	26. 241	10.036	13.578	35.856	51.104	20.505
	SPLEN1000136	63.959	49, 187	29.950	19.924	33.180	31.548	43.178	34.511
	SPLEN1000141	23.876	26, 906	13.071	72.200	12.564	17.177	27.098	35, 500
35	SPLEN1000164	16.339	35.856	18. 324	17.346	11.288	11,589	21.271	23, 356
	SPLEN1000166	24.814	15. 925	15, 170	8. 132	3.719	14. 352	21.337	12.600
	SPLEN1000175	25.901	21.258	13.665	11. 257	7, 394	16.665	16.969	14.377
	SPLEN1000182	18.056	12.663	11.532	12.004	2.626	8. 556	12.618	37, 351
	SPLEN1000185	26.100	41.959	17. 505	17, 472	10.054	14.816	18.440	19.857
	THYMU1000004	44.412	116. 214	81.748	45. 350	91.679	71, 223	84.954	80.324
40	THYMU1000009	92.202	35. 746	24.767	13. 955	26. 373	40.874	48.694	33.357
	THYMU1000015	119.421	76.777	57. 343	70. 294	56. 242	50, 116	65. 925	70.762
	THYMU1000016	74.630	122. 372	55.398	55. 977	36. 943	34, 305	35. 686	44. 484
•				14.380	8. 188	8. 168	17.212	29, 149	11.470
	THYMU1000023	48.992	17. 205			8.770	14.408	18. 502	24.353
	THYMU1000034	23, 593	20. 349	11.577	29.307	3.211	3.832	10.406	7.814
45	THYMU1000035	4.371	10.319	4.870	4.657			15.344	11.760
45	THYMU1000037	20.625	19.668	15. 919	5.609	7.443	10.224 26.145	15.958	
	THYMU1000042	26.144	27.737	22. 945		17.170			21.660
	THYMU1000047	82. 365		47. 962				46.858	56.202
	THYMU1000080	61.757	49. 927	18. 225				51.613	35.091
	THYMU1000094	19.467		53, 131	40.321			33.394	98, 550
	THYMU1000109	149, 316	123.466	67,770	64. 336	47.280	83.698		102, 494
50	THYMU1000127	60.503					36.687	45. 486	19.497
	THYMU1000130	30.806		14. 328					28.159
	THYMU1000137	52.374							22.622
	THYMU1000146	18.567							15,651
	THYMU1000159	70.044	154. 598	47.360	46, 468	39.892	1 00. 239	1 24.033	1,00,001

567

Table 158

	-	110 AFR 11	18.595	78.938	55. 414	75. 224	98. 439	193.301	19. 439
					11.600	12.648	14.819	18.104	21.118
	THYMU1000167	27.217	30.992	12. 427			38. 437	40.004	37.219
	THYMU1000186	98.908	32.919	26.632	32.071	21.873		24. 204	26.183
5	THYRO1000017	32.706	74.720	34.463	29.641	25.692	21.516		25, 420
	THYR01000026	48.577	63.401	19. 205	24. 272	14.810	18. 297	20.045	
	THYR01000034	58.496	36.741	19. 420	21.474	9, 545	24. 247	35. 259	20.763
	THYR01000035	16. 297	9, 507	7.691	4.410	6.774	24. 908	13, 356	9.119
	THYR01000036	24, 463	30.537	12.036	4.674	19.713	10. 204	19.617	16.573
		35.751	45. 426	23.899	33.153	24.644	20.641	51.980	52.710
10	THYR01000040	55, 574	30.112	25. 941	16.711	34.951	38. 423	43.556	36.625
10	THYRO1000061	298. 802	183.339	157.234	94.003	137.727	187.647	208.613	150. 900
	THYR01000067		57.987	43.780	28.114	29,001	66.142	64. 121	39.508
	THYR01000070	129.995	68.453	35. 134	30, 429	26.627	21.975	26.766	30.117
	THYR01000072	48. 939		21.990	11.064	20. 435	19.417	26.995	22.971
	THYR01000084	48. 307	42.611	126.839	102.212	129.747	159.374	206.341	159.771
	THYR01000085	303.121	193.955		6.698	5. 384	7.742	20.711	9. 575
15	THYR01000086	18.728	11.012	7.883		6, 514	3.429	8, 955	5.691
	THYR01000087	13.421	10.853	10.795	1.978	33.042	32.861	31.754	32.557
	THYR01000092	59.642	76.269	32. 514	45.637		20.671	22.009	16.350
	THYR01000093	29. 394	21.625,	13.006	10.368	7.983	25. 517	29.813	20. 577
	THYR01000099	51.966	54.362	21.025	22.941	16. 286		21.662	17. 898
	THYR01000107	29.893	53.294	14.175	20.025	13. 567	9. 104		13, 162
20	THYR01000111	21.644	28.232	19.143	17.545	16.222	11.312	15.745	6. 079
20	THYRO1000121	9.799	13.392	8.363	4.392	6.118	7.651		
	THYR01000124	30.095	17.896	14.782	10.115	12.585	13.557	25.764	13.667
	THYR01000129	30, 388	14.967	9.694	7.939	3. 251	9.857	13.209	9. 668
	THYR01000130	56, 966	72.160	26.934	36.742	15.654	23.842	14.862	25. 040
	THYR01000132	83, 533	105.422	51.451	54. 782	42. 323	42.319	52.428	22, 575
	THYR01000134	33, 349	47.358	20.790	21.807	12.940	21.379	41.470	7. 581
25	THYR01000144	88, 955	17.323	7.936	4.025	4. 431	18.779	29.660	
	THYR01000155	11.674	3, 549	2.761	3.811	1.697	0.000	7.176	2. 825
	THYR01000156	35, 082	28. 027	15. 226	28.722	15.993	22.315	19.772	22. 116
	THYR01000163	68. 114	50.535	54. 325	60.945	50.945	39. 516	29.854	36. 208
	THYR01000173	43,980	34.453	18.714	18,682	5.054	34,676	33.143	20. 988
	THYR01000186	150.529	131.750	70.665	53. 342	44.898	98. 134	69.084	44. 946
30	THYR01000187	89. 162	62.977	42.088	24. 103	13.600	31.751	46. 152	25. 272
	THYR01000190	34,704	43.709	35.680	47.383	21.817	20.074	24.984	29. 176
	THYR01000196	12.960	7.875	6. 426	3.533	5. 208	5.665	10.168	5. 312
	THYR01000197	34.949	40. 382	35.820	20.214	23.273	18. 953	26.665	25. 266
	THYR01000199	19.361	13,983	9.085	8. 320	10.004	7, 851	11.633	10.622
	THYR01000206	47.609	55. 960	31.132	10.479	36.037	22, 453	19.963	14. 483
25	THYR01000221	82.534	81, 160	38. 961	57.909	20.347	30, 565	34. 158	38. 238
35	THYRO1000222	15.768	62, 309	7. 359	7.364	8.966	8.443	11.700	23. 186
	THYR01000228	23.238	16.601	14.212	15.062	17.974	19, 434	9,775	10. 964
	THYR01000241	55.874	49, 255	57.277	39.823	31.045	29.731	25.058	24. 705
	THYR01000242	13.379	26. 177	12.762	19.853	8. 446	8.035	12.464	24. 333
	THYR01000246	7.985	21.129	6. 632	7. 437	5.012	11.050	8.809	26.581
	THYR01000253	60.014	38.765	34, 683	39.349	28. 961	21. 254	21.340	26.307
40	THYR01000270	3,554	0.000	2.696	1.813	2.708	4. 022	4.159	3.250
	THYR01000279	14, 227	10.091	5. 33	3. 542		8. 248	7.649	5.892
	THYR01000285	56.886	54.148	33.944			32. 641	25.655	26.150
	THYR01000288	12, 236	23.331	7.80	4. 959				6.433
	THYR01000296	58.849	34, 305		1 18.781				24.815
	THYR01000320	40, 309			7   13.455				23.604
45	THYR01000322	24.627			2   13. 220	24. 263			17.831
	THYR01000327	26, 339			6.909				12.537
	THYR01000343				4 6.474	9, 696			
	THYRO1000345					4. 038			
	THYR01000358					36, 183			
	THYRO1000358								
50	THYR01000375								
50	THYRO1000381					3.83			
	THYR01000387								
	THYR01000194					8 48.70			
	THYR01000195						3 45. 02	3 48.651	32.418
	11110100133	1 3							

Table 159

			30.000 1	17.7.0	10 000		49 446	(5 000	40 444
	THYR01000400	29. 261	30.808	14,649	12.890	12. 143	17.419	17.865	20. 330
	THYR01000401	48.109	37.938	22.638	16.225	12.893	28. 523	30.627	19. 454
	THYRO1000407	20. 235	11, 480	8.357	3,709	9.881	7, 211	13.275	7.999
5	THYR01000420	58.894	43.095	35, 789	24.115	20. 938	30.086	31.852	26.083
•	THYR01000438	33. 270	20. 145.	29.159	31.273	12.085	10. 585	9. 246	12.932
				27. 337	22.464	17.753	26. 548	22.201	22. 293
	THYR01000452	53.893	37. 152						
	THYR01000455	2. 280	0.834	0.000	0.976	0.585	1. 280	2.641	1.093
	THYR01000471	47. 958	25. 563	18. 216	14.664	13.684	14. 209	21.695	20.773
	THYR01000481	31.917	26. 285	16.526	11,506	15.682	19. 322	21.433	19.881
10	THYR01000484	105. 986	101.654	49.570	65.106	44.560	42.375	53.422	61.358
	THYRO1000488	10.604	11,718	5. 980	2.408	1.075	2. 903	5. 387	5. 572
	THYR01000501	27.472	26, 976	14. 433	9, 731	5. 970	14. 226	13.623	20.839
	THYRO1000502	5. 447	3.089	4. 285	1.572	3.996	4. 353	4. 902	2.744
		4. 701	9. 342	2,729	1,539	2.859	3.412	6.900	4.379
	THYRO1000505			15.352	10.179	15. 441	15.802	32.978	26. 549
	THYRO1000535	36. 284	36.608			21.538	34.069	54.689	
15	THYR01000556	98. 555	26.955	23.471	9. 941				21.009
	THYR01000558	40. 392	34. 267	23.559	20.713	24.798	17, 657	28.862	27.336
	THYR01000569	873.069	308.078	372.545	155.422	299.039	483. 635	445. 882	305. 921
	THYRO1000570	35. 246	19.469	12.612	19.448	7.219	17. 186	18.803	17. 396
	THYR01000572	39.801	10.089	11.294	4.705	3. 845	19.606	13, 915	6.846
	THYR01000573	16.251	10.017	7.249	4.045	4. 497	4.783	12.198	5. 097
	THYR01000577	10.585	9, 999	5.259	3.391	2.076	7. 540	7.747	5,771
20	THYR01000580	39. 072	33.754	20, 407	32.861	20.138	22.441	26, 841	38.662
	THYRO1000584	56.308	33.150	19.548	14.340	19.384	30, 365	39, 545	23. 407
	THYRO1000585	43. 561	24.758	28. 265	16.580	20.169	22. 132	27.817	24. 366
	THYRO1000596	2.673	0.776	0.000	0.000	4.716	2.198	2.119	1.933
					45.120	38. 294	42. 518	37.044	37, 636
	THYR01000602	94. 197	75.969	43. 440			20. 183	23.421	15. 563
05	THYR01000605	37.030	19. 281	9.512	7.831	12.501			
25	THYR01000615	15.039	14.895	6.698	8.884	6.498	7. 491	8.656	8, 551
	THYR01000625	49.869	34. 253	18.419	29.529	18. 526	18. 214	20.134	21.544
	THYR01000636	32. 799	20.827	9, 591	8.974	11.041	15. 919	22.990	18.387
	THYRO1000637	35. 581	23.050	18.908	14, 371	24.139	16.485	41,751	19.963
	THYRO1000641	28. 962	17.660	13.853	8.774	18. 253	16.722	20, 366	17.183
	THYRO1000657	66.685	48.553	43.153	26.769	20.514	33.412	27.427	59.913
30	THYRO1000658	101.090	94.403	57.365	65.686	46.570	42.965	15.054	51.149
	THYRO1000662	30, 501	28.754	7.936	6.202	14.884	24.631	23.740	14. 132
	THYRO1000666	56.263	27.128	11.520	10.878	12.343	19.483	26.494	16.400
	THYRO 1000676	46.904	34.507	12.093	23. 243	14.596	11.035	13.272	18.504
	THYRO1000678	12.599	11.709	10.630	7.426	8. 273	5.498	7,825	12.309
	THYR01000684	61.875	24.579	20. 434	9.128	13.986	27.123	42.335	20.023
	THYRO1000694	94. 566	65,001	36. 187	11.784	39.648	50.883	109.147	47.741
35	THYRO1000699	228. 022	178.345	154. 501	107.031	135. 907	157.164	148.138	139.950
			120.229	65. 349	78.931	61.796	42.847	42.817	59.069
	THYRO1000712	66. 420			12.645	16. 476	20. 968	33, 909	18.460
	THYRO1000715	52. 182	30.514	16.829			8. 581	20.893	12. 979
	THYR01000716	34.776	27.524	13.457	11.085	11.113	21.072	22.860	29. 727
	THYR01000717	64. 920	84.125	21.513	31.324	22.570			
40	THYR01000723	6.184	6.744	4. 434	3.785	5. 307	2.617	6.718	7.719
40	THYR01000734	15.193	18.494	9.892	17.212	6.183	7.960	17.862	10.472
	THYR01000748	94. 224	47.484	24. 348	16. 194	34. 311	34.308	68.067	29.440
	THYRO1000755	24.375	26.453	17.994	18.096	13.613	21, 492	17.967	32.148
	THYRO1000756	50.530	55. 357	19.662	10.236	15. 906	24.457	28,624	19. 162
	THYRO1000776	24.132	29.551	15.488	11.113	9. 272	17.530	17.901	15. 200
	THYR01000777	18.780	26.388	14.190	9.047	9.368	16.446	29.480	15.416
45	THYR01000779	1.795	0.000	0.000	2.494	7.457	0.000	6, 362	2. 532
	THYRO1000782	47, 931	38, 121	28.062	11.863	22.874	28. 629	25, 106	23.954
	THYR01000783	25.655	14.286	12.376	5.578	6. 270	12.787	17, 848	13.045
	THYRO1000786	52.665	48.137	29.971	29.960	23.410	37.344	61.708	40.990
	THYRO1000787	300.022	78.369	95. 279	31.225	58.114	149.896	140, 608	55. 131
	THYRO1000792	56.669	16.981	17.506	14.737	10. 487	12.435	26. 185	19.757
						7.738	18. 245	14.576	9.048
5 <b>0</b>	THYR01000793	21.782	17.626	12.726	12.269				
	THYRO1000795	35.732	43.199	24.656	10.920	12.277	22.001	20.250	17.634
	THYR01000796	23. 496	27.404	20.088	17.955	13. 259	12.893	15. 542	13. 569
	THYR01000798	46.024	29.017	22.439		17.838	27.756	29.891	12.085
	THYRO1000800	51, 341	77.530	54.957	81.739	91.231	44.745	43.380	63.706

Table 160

	THYR01000805	29.203	24.611	12.889	12.552	8.708	24. 185	31. 195	17.746
		116.955	165.320	75.096	94.269	59. 401	58. 491	42. 135	74.481
	THYR01000815								8. 236
	THYR01000829	23.576	12.796	8.360	10.367	5.365	10.395	15. 475	
5	THYR01000835	25. 167	23.644	13.936	14.093	11.798	32.901	18.905	17. 992
•			44, 053	31.047	36.013	19.347	21.091	20.171	23. 430
	THYR01000843	33.508						10. 459	6. 930
	THYRO1000845	18.033	12. 383	7.953	5. 357	8.714	8.050		
	THYR01000852	26.571	15.703	9.149	9.589	4. 965	8.428	10. 204	11.995
		45.596	37.371	20.596	42.732	32, 911	16.694	31.555	30. 260
	THYRO1000855						38, 134	52, 647	49.076
	THYRO1000865	72.472	80.181	43.954	56.430	21.283			
10	THYR01000866	136.754	43.702	88.564	12.275	34.870	89.966	25. 647	39.646
	THYR01000881	484.415	303.533	220.883	156.089	149.161	314. 435	262. 114	229. 042
			28. 931	14. 132	11.237	15.661	21.378	24. 165	10.595
	THYR01000894	65.638				6. 164	7.972	11, 149	13. 327
	THYRO1000895	19.040	17. 281	11.079	9.005				
	THYR01000916	68.849	51.202	36.286	38.745	35.015	21.936	23. 241	21.349
		378.890	211.431	172.873	110.307	168, 147	239.935	221.829	171.250
	THYRO1000917				14.409	20.225	27.710	40.030	15. 229
15	THYRO1000926	74.104	25. 922	17.751					
	THYR01000934	21.900	17.023	11.309	10.688	4.218	13.887	14. 363	11, 574
	THYR01000951	48, 727	35.250	16.046	12.962	18,778	26.338	19. 255	15. 211
			22.838,	17. 193	11.759	7.673	21.372	18.800	19,736
	THYR01000952	34.577				4. 251	13.753	14.833	20.107
	THYR01000956	37.412	15.001	11, 959	8. 197				
	THYR01000960	40.709	23.743	5. 462	12:106	8. 269	13.882	17.580	15. 391
	THYRO1000961	3.619	4.816	1.934	2.829	5. 229	4.913	6.632	5.076
20		31.761	18, 472	14,773	9.113	13.610	18.567	17, 379	12.630
	THYR01000964				28. 185	28.067	36.041	37, 405	44. 344
•	THYR01000971	64.832	44. 237	30.605					
	THYR01000974	107.219	62.723	34. 195	40.953	32.826	39. 260	30.469	42. 586
	THYR01000975	81.132	53. 975	52, 682	49.142	35.144	34.988	44. 912	43.686
			23.344	30.088	11, 305	15.039	29.019	17, 082	16.694
	THYR01000983	44. 267						14.647	19.412
	THYR01000984	43.136	31.868	22.917	23.200	16. 640	18.941		
25	THYR01000988	77.046	58.963	40. 192	43, 118	60. 580	33.078	20.658	30. 028
	THYR01000991	59, 477	49.735	27. 299	24.412	23.236	36.791	41.514	30.530
					16.574	12.745	22.240	23.460	20.374
	THYR01000999	46. 173	27.320	24.436					29, 299
	THYR01001003	45. 343	40.846	34.059	27.728	30.647	22.758	14.074	
	THYRO1001015	105.149	53.043	34.722	25. 220	29.072	70.219	55.045	37. 157
		55.018	27.688	20, 817	19.166	16.243	14.052	10.907	20.419
	THYR01001016					10, 892	16.822	19. 126	15.036
30	THYR01001022	34.560	25.745	16.566	9. 263				
	THYR01001031	79.734	70.269	57. 437	40.146	30.024	20.905	25. 507	25. 466
	THYR01001033	22.581	21.639	10.233	5.613	5. 972	14.479	22.263	14.812
		50. 552	36.895	25. 102	26.692	22.143	17.789	17.845	24.414
	THYRO1001062					26. 587	31.088	36.388	28.011
	THYRO1001063	75, 298	52.927	34.731	26.645				
	THYRO1001071	15. 221	6. 957	5. 949	2.013	6.433	6. 642	7.145	6. 223
	THYR01001080	47.009	39.873	20.480	18. 101	20.162	20.086	35. 494	27.474
35	THYR01001093	66.980	65.072	31.618	33.554	16.112	27.365	31.863	34.516
				12.169	9.015	5. 970	14.570	15.506	13, 653
	THYR01001100	21.067	15. 255				11.510	9.775	8. 902
	THYR01001102	18.746	18.080	6. 257	4.335	1.730			
	THYRO1001104	18.657	25. 635	14.755	25. 137	12,793	22.720	23.958	26.681
	THYRO1001109	15. 251	15. 230	8. 676	4. 654	5. 820	7.397	12.338	9.739
	THYRO1001113						1 10 040	22.426	17.552
40						1 16 170	1 17 945		
		37.344	45. 395	7. 359	6.259	16.170	12.948		
70	THYRO1001120	80.202	35. 430	22.559	15, 448	18.774	31.803	42.346	22.885
40	THYRO1001120	80.202					31.803 24.648	42.346 45.988	22. 885 38. 643
40	THYRO1001120 THYRO1001121	80. 202 52. 621	35. 430 42. 522	22. 559 27. 046	15. 448 29. 236	18.774 28.248	31.803	42.346	22.885
40	THYRO1001120 THYRO1001121 THYRO1001128	80. 202 52. 621 136. 958	35. 430 42. 522 100. 049	22.559 27.045 61.329	15, 448 29, 236 56, 461	18.774 28.248 53.098	31.803 24.648 51.086	42.346 46.988 60.358	22. 885 38. 643 56. 767
40	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001133	80. 202 52. 621 136. 958 94. 452	35. 430 42. 522 100. 049 101. 822	22.559 27.046 61.329 62.367	15, 448 29, 236 56, 461 57, 536	18.774 28.248 53.098 40.128	31.803 24.648 51.086 46.930	42.346 46.988 60.358 37.716	22.885 38.643 55.767 49.125
	THYRO1001120 THYRO1001121 THYRO1001128	80. 202 52. 621 136. 958	35. 430 42. 522 100. 049 101. 822 17. 461	22. 559 27. 045 61. 329 62. 367 8. 019	15, 448 29, 236 56, 461 57, 536 4, 846	18.774 28.248 53.098 40.128 6.568	31.803 24.648 61.086 46.930 9.163	42.346 46.988 60.358 37.716 14.613	22. 885 38. 643 56. 767 49. 125 12. 344
•	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001133 THYRO1001134	80. 202 52. 621 136. 958 94. 452	35. 430 42. 522 100. 049 101. 822	22.559 27.046 61.329 62.367	15, 448 29, 236 56, 461 57, 536	18.774 28.248 53.098 40.128 6.568 2.274	31.803 24.648 61.086 46.930 9.163 4.180	42.346 46.988 60.358 37.716 14.613 3.267	22. 885 38. 643 56. 767 49. 125 12. 344 3. 903
	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001133 THYRO1001134 THYRO10011142	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374	22, 559 27, 046 61, 329 62, 367 8, 019 4, 501	15, 448 29, 236 56, 461 57, 536 4, 846	18.774 28.248 53.098 40.128 6.568 2.274	31.803 24.648 61.086 46.930 9.163	42.346 46.988 60.358 37.716 14.613	22. 885 38. 643 56. 767 49. 125 12. 344
45	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001134 THYRO1001134 THYRO1001142 THYRO1001173	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303	15. 448 29. 236 56. 461 57. 536 4. 846 1. 699 99. 619	18.774 28.248 53.098 40.128 6.568 2.274 143.648	31.803 24.648 61.086 46.930 9.163 4.180 173.339	42.346 46.988 60.358 37.716 14.613 3.267 189.443	22.885 38.643 56.767 49.125 12.344 3.903 126.977
	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001133 THYRO1001142 THYRO1001173 THYRO1001173 THYRO1001173	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 519 6, 214	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354	31.803 24.648 51.086 46.930 9.163 4.180 173.339	42.346 46.988 60.358 37.716 14.613 3.267 189.443 23.098	22.885 38.643 56.767 49.125 12.344 3.903 126.977 12.914
	THYRO1001120 THYRO1001121 THYRO1001123 THYRO1001133 THYRO1001134 THYRO1001142 THYRO1001173 THYRO1001173 THYRO1001177	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556	31.803 24.648 61.086 46.930 9.163 4.180 173.339 14.774 23.552	42.346 46.988 60.358 37.716 14.613 3.267 189.443 23.098 39.234	22.885 38.643 56.767 49.125 12.344 3.903 126.977 12.914 27.932
	THYRO1001120 THYRO1001121 THYRO1001123 THYRO1001133 THYRO1001134 THYRO1001142 THYRO1001173 THYRO1001173 THYRO1001177	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556	31.803 24.648 51.086 46.930 9.163 4.180 173.339	42.346 46.988 60.358 37.716 14.613 3.267 189.443 23.098	22.885 38.643 56.767 49.125 12.344 3.903 126.977 12.914
	THYRO1001120 THYRO1001121 THYRO1001121 THYRO1001133 THYRO1001134 THYRO1001174 THYRO1001173 THYRO1001177 THYRO1001177 THYRO1001177	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781 80, 715	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556 51.976	31.803 24.648 61.086 45.930 9.163 4.180 173.339 14.774 23.552 45.521	42, 346 46, 988 60, 358 37, 716 14, 613 3, 267 189, 443 23, 098 39, 234 44, 962	22. 885 38. 643 55. 767 49. 125 12. 344 3. 903 126. 977 12. 914 27. 932 108. 449
	THYRO1001120 THYRO1001121 THYRO1001121 THYRO1001133 THYRO1001134 THYRO1001134 THYRO1001173 THYRO1001175 THYRO1001175 THYRO1001189 THYRO1001194	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781 80, 715 31, 797	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556 51.976	31.803 24.648 61.086 46.930 9.163 4.180 173.339 14.774 23.552 45.521	42, 346 46, 988 60, 358 37, 716 14, 613 3, 267 189, 443 23, 098 39, 234 44, 962 16, 508	22.885 38.643 56.767 49.125 12.344 27.932 108.449 31.677
	THYRO1001120 THYRO1001121 THYRO1001121 THYRO1001133 THYRO1001134 THYRO1001174 THYRO1001173 THYRO1001177 THYRO1001177 THYRO1001177	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781 80, 715 31, 797 17, 477	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556 51.976 22.784	31.803 24.648 61.086 46.930 9.163 4.180 173.339 14.774 23.552 45.521 16.960 29.010	42, 346 46, 988 60, 358 37, 716 14, 613 3, 267 189, 443 23, 098 39, 234 44, 962 16, 508 29, 959	22.885 38.643 56.767 49.125 12.344 3.903 126.977 12.914 27.932 108.449 31.677 20.054
	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001134 THYRO1001134 THYRO1001173 THYRO1001175 THYRO1001175 THYRO1001189 THYRO1001194 THYRO1001194	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781 80, 715 31, 797 17, 477	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556 51.976 22.784	31.803 24.648 61.086 46.930 9.163 4.180 173.339 14.774 23.552 45.521	42, 346 46, 988 60, 358 37, 716 14, 613 3, 267 189, 443 23, 098 39, 234 44, 962 16, 508 29, 959	22.885 38.643 56.767 49.125 12.344 27.932 108.449 31.677
45	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001134 THYRO1001134 THYRO1001173 THYRO1001177 THYRO1001177 THYRO1001177 THYRO1001189 THYRO1001204 THYRO1001204	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393 444. 098	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084 372. 962	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874 .225. 154	15, 448 29, 236 56, 461 57, 536 4, 846 1, 699 99, 619 6, 214 23, 781 80, 715 31, 797 17, 477 217, 033	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556 51.976 22.784 14.104 189.087	31.803 24.648 61.086 45.930 9.163 4.180 173.339 14.774 23.552 45.521 15.960 29.010	42. 346 46. 988 60. 358 37. 716 14. 613 3. 267 189. 443 23. 098 39. 234 44. 962 16. 508 29. 959 214. 186	22. 885 38. 643 56. 767 49. 125 12. 344 3. 903 126. 977 12. 914 27. 932 108. 449 31. 677 20. 054 193. 594
45	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001134 THYRO1001134 THYRO1001173 THYRO1001175 THYRO1001177 THYRO1001177 THYRO1001189 THYRO1001190 THYRO1001205 THYRO1001205	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393 444. 098 59. 798	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084 372. 962 77. 150	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874 .225. 154 45. 729	15. 448 29. 236 56. 461 57. 536 4. 846 1. 699 99. 619 6. 214 23. 781 80. 715 31. 797 217. 033 51. 526	18.774 28.248 53.098 40.128 6.568 2.274 143.648 10.354 36.556 51.976 22.784 14.104 189.087	31.803 24.648 61.086 45.930 9.163 4.180 173.339 14.774 23.552 45.521 16.960 29.010 245.605 26.773	42. 346 46. 988 60. 358 37. 716 14. 613 3. 267 189. 443 23. 098 39. 234 44. 962 16. 508 29. 959 214. 186 26. 362	22. 885 38. 643 56. 767 49. 125 12. 344 3. 903 126. 977 12. 914 27. 932 108. 449 31. 677 20. 054 193. 594 35. 040
45	THYRO1001120 THYRO1001121 THYRO1001121 THYRO1001133 THYRO1001134 THYRO1001173 THYRO1001173 THYRO1001177 THYRO1001177 THYRO1001194 THYRO1001104 THYRO1001204 THYRO1001204 THYRO1001204 THYRO1001213	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393 444. 098	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084 372. 962 77. 150 51. 273	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874 225. 154 45. 729 33. 830	15. 448 29. 236 56. 461 57. 536 4. 846 1. 699 99. 619 6. 214 23. 781 80. 715 31. 797 17. 477 217. 033 51. 526 51. 454	18.774 28.248 53.098 40.128 40.568 2.274 143.648 10.354 36.556 51.976 22.784 14.104 189.087	31.803 24.648 61.086 46.930 9.163 4.180 173.339 14.774 23.552 45.521 15.960 29.010 245.605 26.773 34.214	42. 346 46. 988 60. 358 37. 716 14. 613 3. 267 189. 443 23. 098 39. 234 44. 962 16. 508 29. 959 214. 186 26. 362 24. 649	22. 885 38. 643 56. 767 49. 125 12. 344 3. 903 126. 977 12. 914 27. 932 108. 449 31. 677 20. 054 193. 594 47. 409
45	THYRO1001120 THYRO1001121 THYRO1001121 THYRO1001133 THYRO1001134 THYRO1001173 THYRO1001173 THYRO1001177 THYRO1001177 THYRO1001194 THYRO1001104 THYRO1001204 THYRO1001204 THYRO1001204 THYRO1001213	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393 444. 098 59. 798 53. 123	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084 372. 962 77. 150 51. 273	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874 225. 154 45. 729 33. 830	15. 448 29. 236 56. 461 57. 536 4. 846 1. 699 99. 619 6. 214 23. 781 80. 715 31. 797 17. 477 217. 033 51. 526 51. 454	18.774 28.248 53.098 40.128 40.568 2.274 143.648 10.354 36.556 51.976 22.784 14.104 189.087	31.803 24.648 61.086 46.930 9.163 4.180 173.339 14.774 23.552 45.521 15.960 29.010 245.605 26.773 34.214	42. 346 46. 988 60. 358 37. 716 14. 613 3. 267 169. 443 23. 098 39. 234 44. 962 16. 508 29. 959 214. 186 26. 362 24. 649 49. 913	22. 885 38. 643 55. 767 49. 125 12. 344 3. 903 126. 977 12. 914 27. 932 108. 449 31. 677 20. 054 193. 594 47. 409 35. 697
45	THYRO1001120 THYRO1001121 THYRO1001128 THYRO1001134 THYRO1001134 THYRO1001173 THYRO1001175 THYRO1001177 THYRO1001177 THYRO1001189 THYRO1001190 THYRO1001205 THYRO1001205	80. 202 52. 621 136. 958 94. 452 17. 941 10. 016 315. 863 38. 323 65. 825 71. 764 43. 753 24. 393 444. 098 59. 798	35. 430 42. 522 100. 049 101. 822 17. 461 5. 374 215. 361 13. 237 73. 170 109. 416 58. 316 20. 084 372. 962 77. 150	22. 559 27. 046 61. 329 62. 367 8. 019 4. 501 158. 303 7. 198 30. 535 54. 067 68. 460 15. 874 .225. 154 .45. 729 33. 830 27. 897	15. 448 29. 236 56. 461 57. 536 1. 699 99. 619 6. 214 23. 781 80. 715 31. 797 17. 477 217. 033 51. 526 51. 454 20. 382	18, 774 28, 248 53, 098 40, 128 6, 568 2, 274 143, 648 10, 354 36, 556 51, 976 22, 784 14, 104 189, 087 31, 541 44, 844 32, 686	31.803 24.648 51.086 46.930 9.163 4.180 173.339 14.774 23.552 45.521 16.960 29.010 246.605 26.773 34.214 50.109	42. 346 46. 988 60. 358 37. 716 14. 613 3. 267 189. 443 23. 098 39. 234 44. 962 16. 508 29. 959 214. 186 26. 362 24. 649 49. 913	22. 885 38. 643 56. 767 49. 125 12. 344 3. 903 126. 977 12. 914 27. 932 108. 449 31. 677 20. 054 193. 594 47. 409

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Table 161

	-	717 300 3	T	12 000 1	40 449 1	30 406	72 140	00 614 7	50. 152
	THYR01001258	115. 192	68. 322	37.962	28. 447	39. 496	73.140	89. 614	
	THYR01001262	29. 592	38. 992	24.922	22.829	20. 191	14. 595	14. 682	19. 309
	THYRO1001266	24. 695	23.851	16.014	7.968	11.573	18. 488	19.268	13, 434
5	THYR01001271	37.090	37.276	12.145	11.215	2.868	19. 505	17. 992	11.460
	THYR01001287	69. 292	40.644	17.033	16.333	18. 990	24. 523	40. 591	27.350
	THYR01001290	38. 183	9.778	9. 132	6.909	7.883	17. 550	22.844	10.046
	THYRO1001291	27.456	31, 200	13.335	8.894	13.643	16. 343	24. 246	14.305
	THYR01001297	22.802	40, 193	15.454	24. 356	18.908	13. 849	19.636	27.811
	THYR01001302	32.724	25.039	21.076	11.586	19. 524	23.410	57.069	22. 259
10	THYR01001313	54, 483	44.710	22.791	17.196	22.860	28. 535	38.530	26.619
	THYR01001320	67. 151	79.399	38.582	43.377	31.441	31.488	30.487	34. 150
	THYR01001321	32. 185	46.760	20.156	31, 133	26.936	21.803	17.729	26. 264
	THYR01001322	56.040	44.139	25. 288	32.717	28.245	19.900	28.415	30.093
	THYR01001327	11. 598	12.117	3.614	3, 130	6.285	5.136	8. 978	9. 997
	THYR01001336	45. 342	100.054	38, 339	43.663	34.416	23.794	31. 249	61.226
4.5	THYR01001347	8, 316	11.569	4.451	4. 135	3.827	2.861	6.260	3. 931
15	THYR01001358	96.749	91.718	27.513	38.148	45.764	39, 905	54. 447	48. 267
	THYR01001363	76. 229	50.596	45.707	32.563	22.003	40.930	35. 965	23.714
	THYRO1001365	63.340	44.755,	24.569	15. 278	14, 500	31, 255	62.023	22. 216
	THYRO1001374	80.359	54.703	28.941	21.895	26.409	86.809	59.724	67. 154
	THYR01001401	138. 528	81.793	116: 025	115.772	62.059	81.850	81.710	84. 369
	THYR01001403	75.077	60. 253	47.159	43.576	31, 391	38,040	41.579	34. 801
20	THYR01001405	75.788	63. 929	37.018	66.708	25.398	44. 268	169.777	92. 288
	THYRO1001406	99.789	119.681	106.617	111.553	73. 294	82.322	63.741	106, 694
	THYRO1001411	164. 801	155. 374	122.876	101.166	90.616	97.554	90, 344	81, 141
	THYR01001420	467.850	125. 400	141.742	95, 133	79.850	256.705	243.974	168. 095
	THYR01001426	179.694	226.744	136.659	182.920	57.912	158.699	76.886	79. 382
	THYR01001430	42. 233	36.308	24. 265	13.334	24. 942	28. 220	31.096	31.763
25	THYRO1001434	109. 844	40. 429	23. 142	7.076	19.838	16.721	46.971	13. 594
	THYR01001456	86.810	51.093	28. 872	22.686	29. 334	38.972	42.073	32. 789
	THYRO1001457	98. 410	46.954	51. 922	44. 428	26. 365	58.702	73, 800	71. 948
	THYRO1001458	142, 203	61.648	63.756	91.611	29. 372	63.294	57. 491	83.860
	THYR01001459	98. 569	70.732	48. 940	49.572	33.394	53.365	59. 458	61. 428
	THYR01001471	29.011	30.922	22.501	12.339	12. 979	11.855	19.026	15.004
30	THYR01001478	88.744	24.933	23.684	23.261	16.773	41.417	28.941	16.857
30	THYR01001480	198, 549	217. 139	159.064	171.096	130.028	151.021	98.977	203.804
	THYR01001481	72.983	76. 982	51.877	37.940	41, 871	34. 156	32.190	31.811
	THYR01001487	156, 213	112.142	110. 985	77.310	74.839	88.309	62.208	64.884
	THYRO1001495	60, 311	64. 175	75. 269	57.588	39. 964	22.882	51.168	78.626
	THYR01001498	60. 093	50. 240	28. 962	43.623	28.080	27.349	44. 121	57.310
	THYR01001510	78. 106	71.131	37.969	22.613	29. 598	45. 141	25.613	34,714
35	THYRO1001512	146. 930	95. 726	82, 300	47. 386	70.311	138.360	105.274	87.137
	THYR01001519	143. 411	115. 340	57.861	92.182	36.860	89. 655	54.540	72.487
•	THYR01001522	86. 178	52.213	40. 302	33.014	28.267	48.497	38. 421	32.647
	THYR01001523	42.807	21.996	19.646	7.023	13.176	31.304	17.358	26.586
	THYR01001526	28.272	36.470	18, 141	20.984	18. 220	25.059	22.056	22. 382
	THYR01001529	56. 422	40.050	50.636	49. 921	36, 172	38.431	43.929	41.984
40	THYR01001534	79.983	41.665	36.130	45.070	31.736	27.199	39.647	22.708
	THYRO1001537	266.845	336, 357	127. 186	167, 167	121.366	235. 919	269.119	105. 552
	THYR01001541	184. 924	142.434	89.266	94.007	73, 101	77.708	42.435	36.282
	THYR01001545	45, 721	28.807	17.637	23.355	11.596	33.223	26.025	32,640
	THYRO1001559	30. 285	28.050	27.503	21.583	24.440	18.855	21.731	20.280
	THYR01001563	81.147	53, 590	40.132	34.989	31.762	54.315	46.120	51.808
45	THYR01001570	160, 698	53. 241	43.074	13, 542	48, 479	91.833	66.191	35.765
_	THYRO1001573	121. 318	40.895	58.993		41.403	54.710	52, 876	28.623
	THYR01001584	69. 312	78. 135	36.886	44, 973	43.785	43.480	40.786	52.141
	THYR01001593	44. 626	47. 299	8.544	35.805	8.587	5.747	5.738	4.447
	THYR01001595	86.656	81. 363	41.727	44. 260	36, 433	28.946	28.668	31.638
	THYR01001596	68, 810	32. 126	33.747	19.824	25. 437	33.051	41.347	20.355
50	THYR01001602	83.486	75.627	45. 307	63.834	30.332	45.771	44.672	49.010
50	THYR01001605	44. 557	32.876	26.697			5. 627	17.556	16.676
	THYR01001608	155. 484	67. 359	43. 850		31.843	58. 215	58.920	39.494
	THYR01001617	84. 352	72.661	68, 377	48. 198	54.691	37. 282	30.097	54.744
	THYR01001634	61.852		24. 126		22.530	35. 972	34.077	28.775
	101001001034	01.032	33.133	1 64.128	1 10.041	1	1		

Table 162

	(W. E. & A. D. & A. & A		126 666 1	100 651 19	17 004	62 677 1	74. 963	57.380	70.174
					17.804	62.577		31.301	36.440
	THYR01001641	56.288	37. 515	23. 987	22.669	22.635	47. 096 20. 671	23.336	16.130
	THYR01001656	46.272	34.075	24. 272	14, 259	16. 135	23. 345	31. 267	21.631
5	THYR01001658	38.715	35, 384	15. 215	12.669 15.015	9, 537	9.794	20.777	17, 147
	THYROTOOTEGT	12.296	22.714	50.424	34. 364	50.747	38. 082	34.858	41,054
	THYR01001671	50.011	59. 547		17. 925	41. 381	103. 416	95. 249	37.062
	THYR01001672	174.047	48.626	52.990 41.886	44. 045	40. 533	34.065	30. 562	33.114
	THYRO1001673	84.547	78. 591	53. 250	75. 184	37. 282	129.575	50. 337	112,501
	THYR01001677	115. 789	184, 195	56.368	28. 898	58, 930	62. 855	51. 341	29. 701
10	THYR01001683	38.015	42. 900		16.461	15. 124	50.006	58. 501	25.463
	THYR01001700	96.033	45. 482	30. 258	45. 679	27.558	56. 203	56.767	54. 824
	THYR01001702	104. 525	90.670	66, 901 65, 413	39. 114	40. 388	88.732	101.241	68. 988
	THYR01001703	130.645	112.852	58. 522	50.870	37.126	36.387	37.277	63.203
	THYR01001706	91.082	82.049	20. 543	5. 921	22. 162	9. 493	31.475	17.215
	THYR01001721	34. 852	21.558		28. 557	23. 707	34. 262	31.779	30.693
15	THYR01001725	49.609	39. 621	22.513 161.719	64. 173	142, 140	284. 093	229, 429	104, 416
	THYR01001710	401, 603	145. 337	33.629	38.777	22.430	45. 582	54. 154	54, 913
	THYR01001738	89. 896	75. 892	27.130	12.056	9. 553	33, 154	29.680	20.832
	THYR01001743	49. 231	21.758, 17.745	12.052	5.744	9. 946	20.567	17, 357	15, 138
	THYRO1001745	34.753	37.766	23. 996	18.634	16, 249	24.636	33.799	27.306
	THYR01001746 THYR01001770	41.622 103.357	62.531	51.786	43.073	39. 785	65. 980	54. 332	47, 446
20	THYRO1001772	129. 127	129, 155	79.515	82.371	76.101	53.649	49, 368	77.136
	THYR01001778	384. 882	146.526	97.702	61. 349	90.096	136.302	175. 998	85.468
	THYRO1001778	105.591	94.089	51.614	51.310	47.627	57.471	55. 262	69.224
	THYR01001796	218.755	90.413	86.089	46.396	63.339	153.810	148.699	63.431
	THYR01001800	89. 126	64. 948	37.534	20.212	33. 235	41.405	36.130	25.761
	THYRO1001803	272.135	195, 625	179.931	121.130	156, 151	183.032	218.545	154.914
25	THYR01001809	58. 170	31.728	28.593	29, 699	25.633	36.954	29.839	25, 467
23	THYR01001817	64.728	50.418	26.089	15. 924	19.828	34. 567	51.140	43.878
	THYR01001819	190. 982	76.509	54. 579	22.923	63.162	79.239	96.822	48.339
	THYR01001828	234.551	180.238	92. 244	80.148	104.168	85.912	160. 310	122, 500
	THYR01001854	219.242	211.323	112.250	150.918	95.727	100.608	75. 437	109, 696
	THYR01001895	44.632	35.971	20.836	14. 220	19.503	17.351	23. 442	22.241
30	THYR01001907	93.660	85.352	41.680	44.441	40.868	38.888	56. 595	43.717
30	TRACH1000006	33.077	27.517	13.610	11.659	11.195	23.390	21.396	13.682
	TRACH1000013	26.029	19.365	8.037	11.958	5.076	14. 402	20.496	12.167
	TRACH1000074	86.302	70.850	32.892	34. 317	28, 366	44.067	58. 165	52.228
	TRACH1000095	48.021	44, 110	17.672	16.895	20.410	35.389	47. 442	40.607
	TRACH1000102	160, 667	128.745	55. 282	64. 147	57. 430	67.455	96.519	73.638
35	TRACH1000108	25. 597	37.670	13. 402	14.907	16.504	16.136	17.158	22.858
33	TRACH1000126	77.681	74.516	36.350	26.803	33.821	49.762	65, 600	50.277
	TRACH1000145	73.548	74.493	25. 762	17.947	22.979	32.054	38. 447 33. 858	1 25.115
	TRACH1000160	48.076	58. 220	20.043	15.138	20.069	33.175	53.846	52.975
	TRACH1000184	91.686	86.638	74. 932	279.361	88. 220	48.252 26.122	42.606	29.900
	VESEN1000004	62.054	51.690	18. 581	21.964	17. 610 27. 518	43.145	68.086	49, 423
	VESEN1000007	99, 131	44.516	29.577	21.187	35. 545	51.737	104.132	45. 994
40	VESEN1000013	171.250	57.002	99, 295		64. 165	105. 318	97. 599	79.474
	VESEN1000028	154.863	100. 292			39, 182	86.619	98.065	63.442
	VESEN1000059	144. 402	97.274	74. 579		34, 794	116.439	77. 465	65.031
	VESEN1000100	189, 864	54,735	41.418		31.521	66.087	66.041	39.378
	VESEN1000107	86. 037 75. 673	47, 432	28. 526		24.766	41.803	57. 926	29.302
	VESEN1000117	58.990	42.673	43. 051		31.414	56.718	73.188	51.792
45	VESEN1000122 VESEN1000137	28.827	12.637	7.708			19,000	24. 465	12.213
		163. 283	63.672	50. 465			54.086	101.701	58.407
	VESEN1000195	9.881	2.089	6.413					
		402,741	271.057						
	VESEN1000279	302, 568	148.812	122.811			148.698		
	VESEN1000363 VESEN1000388	162. 477							
50	VESEN1000394	142. 530							
	VESEN1000410	136. 126							
	VESEN1000411	95. 259							59.117
	VESEN1000415	97. 225							
	IESENTUUU415	1 31.223	1 23.333	1	. 1 30.540				

Table 163

	VESEN1000440	101.690	47,149	49.195	32.607	27.881	49. 154	45. 485	40, 340
	VESEN1000452	188. 242	75.844	67.861	21.929	49.688	101.557	105.023	55.625
	VESEN1000539	393.622	128, 413		155. 268	285.073	217.892	156.970	105.498
_			40, 448	28. 459	17. 920	17. 204	20.338	40. 271	30.185
5	VESEN1000554	44.150	50. 564	47. 257	21.505	36.349	59.158	68.956	34.611
	VESEN1000557	108.763				41.610	59.636	65, 502	37.895
	YESEN1000575	151. 228	53.084	39. 503	26.612			80.130	52, 937
	VESEN1000585	106. 127	43.069	41.516	30.022	40.857	51.129		
	VESEN1000592	3.732	4. 371	1.727	2.763	2.784	4. 336	0.000	0.000
	VESEN1000658	122.632	54.799	53.689	27.783	41.778	66, 943	69.146	46.823
10	VESEN1000669	454. 284	184.969	184.094	116.303	152.848	275, 995	209.035	150.917
· -	VESEN1000743	93. 271	55.577	38.667	37.030	25. 203	47.385	47.073	46.048
	VESEN 1000752	132.397	105.539	71.129	71.113	87.050	96.768	63.315	77, 177
	VESEN1000761	58.860	37.210	39. 232	28.055	41.286	48.665	37.844	25. 644
	VESEN2000039	1610.708	423. 257	575, 130	281.845	514.008	1029.335	742.044	261.643
	VESEN2000102	157.000	68.371	47.526	31.817	43.456	78.881	87.904	46.756
	VESEN2000164	67.615	99.316	47.555	50. 732	57.545	101, 472	141.913	60.455
15	VESEN2000175	11.198	3.920	4.227	2.329	1.448	2.820	3, 186	3,710
	VESEN2000186	302.893	166.977	128.067	101. 481	89.845	151.983	136.632	157.737
	YESEN2000199	364.016	262.765,	186.502	152.072	152.565	198.826	191.332	195.185
	VESEN2000200	61.361	28.617	25.760	13.454	12.471	25.754	39.784	31.121
		59.937	29, 170	19.088	10.312	16.203	30.641	61.987	24. 109
	VESEN2000204		34. 497	30. 351	21. 300	16.675	31.656	29.879	27.886
20	VESEN2000218	46.156			30.651	31.117	44. 365	42.098	43.558
	VESEN2000230	87.277	57. 160	38.252			15, 146	23. 503	20.851
	VESEN2000272	18, 326	25.046	19. 526	14.701	21.471		38. 504	23.627
	VESEN2000299	81.003	29.068	28.969	16.886	22.798	37.073	52.687	
	VESEN2000323	102. 974	73.231	65.632	62.476	64.170	44.083		53.681
	YESEN2000327	273.358	190.493	102.117	60. 523	95.669	114, 144	160.249	65. 341
	VESEN2000128	52.003	27. 894	15.775	9.884	11.945	24. 112	26.254	20. 997
25	VESEN2000330	109.315	77.876	36. 193	27. 267	44. 428	48. 237	51.597	44. 132
	VESEN2000336	55.020	22, 112	15.818	14.036	11.558	21.687	27.119	27.342
	VESEN2000354	157.246	74.852	37.950	19. 235	42. 182	51, 559	45. 485	29. 194
	VESEN2000378	66.998	66.140	23.647	15.673	16.217	28.709	41.497	35.393
	VESEN2000379	54.007	68.263	27.636	45. 302	17.881	35. 928	44.060	55. 125
	VESEN2000397	27.834	20.615	10.624	8.727	4.818	15. 386	21.163	18.688
30	VESEN2000416	32.241	18.712	9. 825	8.843	5.474	12.685	17.453	16.485
	VESEN2000420	26.334	9,499	7.013	2.363	5, 104	8. 281	2.634	1.015
	VESEN2000430	18.312	20.459	12.183	7. 101	4.975	13.810	17.050	19.805
	VESEN2000448	39.040	15.163	13.638	4.769	9.693	14.334	26.387	13.923
	VESEN2000449	130.475	60.437	47.055	28.198	46.878	64.756	79.761	49.783
	VESEN2000458	54, 149	49.676	24.294	20. 921	18.957	24,771	39.745	38.640
	VESEN2000562	96. 176	59, 785	49.030	22. 452	26.435	54.420	70.890	48.405
35	VESEN2000573	9, 605	2.326	1.730	0, 480	0.850	3.785	3.113	2.414
	VESEN2000604	89.021	25. 246	24.495	10.300	14.725	40.448	47.664	24.052
	VESEN2000614	309.658	310.143	158.396	121.428	98.306	193.176	285. 544	193, 901
	VESEN2000638	20, 825	13.750	9.472	3.518	5.018	8.616	15. 565	14, 138
	VESEN2000641	48.159	26.214	12.211	7.625	12.728	19.489	34. 963	19.847
	VESEN2000645	59. 209	24. 195	14.955	7.186	18.507	28.178	34. 263	17.733
40	Y79AA1000013	157.258	82.237	47.630	29.858	46. 920	77.296	68, 488	40.042
	Y79AA1000030	243. 192	141.007	106.937	74.649	80.890	166.613	137, 379	98.647
	Y79AA1000033	49, 439	83.718	30.433	22. 365	22.376	31.534	35.936	31.220
	Y79AA1000037	41.732	23.568	14, 154	16.224	15.348	13.136	21.199	16.632
		32.341	27.270	14. 230	18.610	9.838	21.052	18.336	19.147
	Y79AA1000041				58.738	48.608	73. 595	69.769	54.893
	Y79AA1000059	153.140	85.760	57.915		53.004	14, 961	23, 027	24.640
45	Y79AA1000065	29.024	32.383	43.083	35. 688		128. 427	148.052	120.067
	Y79AA1000081	173.505	497.689	138.675	253.938	133.917	<del></del>		1 1 1 1 1
	Y79AA1000127	103.173	80, 281	69.484		62.524	80.674	36.808	16.356
	Y79AA1000130	69.801	86.217	30.612		38.125	34, 801	24. 913	36.307
	Y79AA1000131		1161.128	226.879		215. 457	854. 176	483, 175	1147.374
	Y79AA1000134	127.126	50.652			39.721	89.186	71.223	41.628
50	Y79AA1000143	38.064	56, 092			43. 450	26.885	22.084	33.064
	Y79AA1000144	20.785					12.606		17. 382
	Y79AA1000150	70.908							45. 702
	Y79AA1000153	473.493							511.885
	Y79AA1000155	75.693	64.809	33, 184	43.509	28.975	33.094	32.512	48. 232

Table 164

							26 147 1	45 613	£2 £2£ }
	Y79AA1000179	86.164	94. 446	49.366	55. 296	58.791	60. 147	47.613	52. 526
	Y79AA1000181	80.781	67.215	32. 483	29. 549	31,500	35. 404	49. 327	33.105
	Y79AA1000202	306.822	216.805	147.425	103. 426	137.718	171.835	204. 385	165.929
-	Y79AA1000207	105.429	123.961	53. 587	73.034	70.343	43.214	45, 289	43.628
5			209. 292		183.832	147.889	192.552	228. 518	129. 266
	Y79AA1000214	383.142		15, 620	61.846	15.816	19.929	16.837	13.933
	Y79AA1000222	22.954	21.555			43.039		132.959	53.641
	Y79AA1000226	132.385	77.693	43.017	23. 388				75.560
	Y79AA1000227	115.766	115. 677	67.073	61.011	55.085	60.149	67.002	
	Y79AA1000230	45.896	40.474	17.716	13.218	15.881	19.738	28. 415	24.012
	Y79AA1000231	89.296	107, 825	54.778	82.032	51.998	45.803	58. 529	93. 293
10		50.494	47.587	29.697	28. 552	35.135	32.631	62.953	29.843
	Y79AA1000239		53.770	28. 305	21.170	27.158	24.070	38. 214	28.393
	Y79AA1000258	45.676			30.031	40.259	65. 427	65. 925	48.582
	Y79AA1000268	116.499	61.766	35. 684		26.984	89. 369	74. 183	28. 252
	Y79AA1000269	36.988	41.536	21.854	18. 345				
	Y79AA1000270	70.349	65.424	33.771	24. 490	36.913	20. 903	38.891	30.089
	Y79AA1000280	52, 901	53. 162	47. 984	50.005	29.457	37.014	42. 331	53. 345
15	Y79AA1000285	37, 272	42.207	25, 179	14.304	14, 335	34.8D1	24.865	20. 291
			10.594	11.909	7.559	12.902	8,040	7, 052	8.641
	Y79AA1000295	10.340			64.592	36.178	56. 127	56. 928	58.020
	Y79AA1000307	67.533	64.757,	61.969			94. 564	146. 279	83.495
	Y79AA1000313	224. 230	107.870	95. 224	65.861	65, 836			43.590
	Y79AA10B0314	150.954	88.811	114, 139	31.101	138.725	106. 102	94. 884	
	Y79AA1000328	25.270	21.003	21.314	15. 992	13.358	17.078	25. 728	22.062
20	Y79AA1000334	70.086	48.685	34.036	32.394	26. 966	25. 485	44. 339	35.712
	Y79AA1000342	445. 189	140.661	207.068	102.538	170.033	280.562	201.342	123. 827
	Y79AA1000346	44. 966	28.105	25.613	13.811	29.974	38.613	18.724	15. 227
		183.577	87.476	90.030	89.865	36.284	83.081	92.665	49. 209
	Y79AA1000347			102.606	92.069	66.005	137. 226	121.401	90. 266
	Y79AA1000349	180. 947	135.094			41.342	46. 383	35. 944	35.848
	Y79AA1000355	81.202	61.139	54.018	52.567		35, 299	37. 961	39.102
25	Y79AA1000368	45.079	38. 521	25.612	35. 417	24.877			
	Y79AA1000388	34.856	29.318	53.178	46.283	64.992	15, 602	20.395	27.793
	Y79AA1000392	274.040	169.752	96.625	109.904	62.391	137.141	143.707	98.881
	Y79AA1000405	52.788	38.000	27.665	15. 987	21.983	34.628	36.536	24.328
	Y79AA1000410	367.438	401.406	216.699	294.500	169.645	216.009	99.999	119.786
		19, 321	19, 430	17.167	18.384	13.307	17.286	11.353	16.663
	Y79AA1000420			38.233	39.006	35, 194	25.311	19. 482	25.935
30	Y79AA1000423	54. 384	64. 128			18. 993	28.308	30, 801	19.059
00	Y79AA1000426	51, 920	32.060	27.489	16.208		13.924	19.534	15.486
	Y79AA1000432	31.920	23.564	18. 505	7.033	17.684			
	Y79AA1000453	100.064	106.207	64. 195	87.842	32.741	36.705	43. 951	75.421
	Y79AA1000465	32.600	20.760	8.375	9.114	6.582	11.349	19.307	16.375
	Y79AA1000469	97.006	89.211	57.415	39.971	51.138	78.959	69.898	46.327
	Y79AA1000480	49. 123	43.661	36.763	32.840	25.674	27.684	32.111	29. 981
35		29.200	23.820			29.500	19.819	9, 990	17.119
33	Y79AA1000502					44. 981	81.691	94.837	59.780
	Y79AA1000521	165.752				27.637	22.639	17.299	34.366
	Y79AA1000534						55.445	40.270	39.870
	Y79AA1000538	90.033				53.051		67.204	89.085
	Y79AA1000539						49. 286		
	Y79AA1000540						64.022	69.258	38.304
40	Y79AA1000560		217.439	285.257			163. 480	137.130	150.237
40	Y79AA1000574					16.138	19.256	27.792	16.219
	Y79AA1000584						7.940	8. 373	2.978
							89.842	105.272	87.142
	Y79AA1000589						26.476	39, 963	26, 495
	Y79AA1000598						48.490	27, 953	19.342
	Y79AA1000600							46.377	40. 338
45	Y79AA100060								
45	Y79AA1000611	125.086	117. 263	62.98	3 91.66	44, 430	82.703	59.073	106.707
	Y79AA100062			33.26	3 16.064	26.240			
	Y79AA100063					38. 373			
	Y79AA100064						22.853	24.218	28.136
	Y79AA100065								
50	Y79AA100067								
50	Y79AA100067								
	Y79AA100067								
	Y79AA100068	2 206.91	1 109. 20						
	Y79AA100068			5 30.76	4 23.66	1 15.359	27.974	25.066	30.575
	1137313000	,,,,,,,							

Table 165

	V70441A00E07	593.441	140. 294	205. 250	28.388	180.538	358.317	185.955	157.146
	Y79AA1000697		45. 357	16, 113	12, 299	6,003	17.423	23. 401	24. 353
	Y79AA1000700	21.077	42. 446	9. 035	13.744	21.360	47.616	22.905	32.458
	Y79AA1000702	62.438	7. 058	5. 353	3. 179	5. 193	12.141	10. 206	6.710
5	Y79AA1000704	19.430	17. 592.	10. 298	10,719	8.004	6.779	14. 333	13.157
	Y79AA1000705	10.998		27, 106	19. 428	22.464	33.577	31.373	29.033
	Y79AA1000717	81.752	30.031 18.986	16. 192	21. 995	16. 249	16.693	21.786	16.366
	Y79AA1000722	36.212		22.178	41.307	9. 368	15.888	23.618	33.068
	Y79AA1000724	38. 197	38. 149		20.692	45.339	70.264	50.747	27. 206
	Y79AA1000726	145.871	38. 218	60. 209	19.757	17.790	29. 363	24. 308	23.170
10	Y79AA1000734	39.812	31.718	23.656		5. 845	15.311	22.833	15. 914
	Y79AX1000748	27.090	25. 462	9. 232	12.141	55. 324	60.996	55. 270	67.659
	Y79AA1000750	117. 327	94. 348	68. 179	74.017	0.825	1.965	2.636	3, 173
	Y79AA1000752	1.118	1.818	0. 920	1. 289	18.788	15. 530	30. 498	24.069
	Y79AA1000774	28. 946	29. 201	13.619	12.927	27.128	25. 527	51, 928	36.475
	779AA1000776	62.397	39. 548	26. 589	27.629	19.771	37. 889	47.008	33.768
15	Y79AA1000777	88. D93	76. 872	25. 449	23. 295	21.743	40. 267	35.715	34.927
	Y79AA1000778	89.017	55. 709	39. 247	33. 579	12. 105	36.578	36.040	18. 916
	Y79AA1000782	67.565	23.947	20.966	11.489 20.358	20.827	23.886	24. 033	27.859
	Y79AA1000784	39.988	33.246	27. 325 15. 477	16.092	14. 809	22.064	28. 950	21.010
	Y79AA1000794	41.650	24. 812 ° 25. 329	17. 225	7. 394	11.113	22.848	25. 673	22.742
	Y79AA1000800	41.806 11.595	15.878	4. 838	4. 573	8. 562	8.929	13.772	8.772
20	Y79AA1000802	65.610	45. 406	23.562	18. 162	27.677	27.293	47.887	38. 440
20	Y79AA1000805	63.932	47.479	31.983	34. 426	26.716	43.371	35.784	35. 139
	Y79AA1000823	22.185	48. 954	19. 279	19. 138	20.407	22.530	21.540	22.820
	Y79AA1000824	27.742	25.712	19.443	10. 124	16.886	17.840	25. 211	16.052
	Y79AA1000827	25.479	15. 274	10.916	8.356	10.528	8.349	18. 396	16.070
	Y79AA1000831	72.020	40. 592	97. 281	14. 517	90.381	82.278	84. 325	35. 373
25	Y79AA1000833	471.030	168.358	184. 092	104. 334	176.646	249.032	310, 721	135, 495
23	Y79AA1000850	68.647	36. 187	20.372	16.113	21.247	21.299	56, 582	51.148
	Y79AA1000856	77.469	45.416	31.674	22.522	37.097	33.815	62.486	52.013
	Y79AA1000862	113.504	90.763	34.743	41.876	44. 348	44.281	54.080	52. 382
	Y79AA1000876	9.498	19.259	12.167	B. 739	10.542	5.725	6, 252	6.011
	Y79AA1000888	44.286	18.430	12.128	10.726	16.431	17.727	35.647	22.169
30	Y79AA1000902	25.675	20.186	13.114	21.076	13. 224	15.117	12. 128	12.728
••	Y79AA1000935	349.462	152.766	266.451	85. 379	264. 556	178.067	253.603	154. 565
	Y79AA1000959	32. 431	15. 556	16.803	4.758	23. 529	16.748	16.620	10.584
	Y79AA1000962	37.877	67. 978	25. 428	20. 228	38. 757	20.056	35.087 60.514	28, 250 78, 400
	Y79AA1000963	77. 792	69.690	30.704	66.559	22.376	45. 923	77.798	55.788
	Y79AA1000966	60.459	53.027	38. 303	43. 259	53.012 74.280	58.436 67.804	71.776	42.966
35	Y79AA1000967	112.210	96. 985	52.461	31.773	52. 133	37.934	58.710	32.052
••	8360001VV	57. 156	75.011	31.312 29.787	31.786 20.498	30. 555	33.354	44.510	23.718
	Y79AA1000969	73.694 19.416	47.137 22.033	12. 239	12.727	10.894	13.904	19. 193	13.612
	Y79AA1000976 Y79AA1000978	50. 835	57. 439	51.253	31.538	53.350	33.330	50.341	51.246
	Y79AA1000918	162. 170	116.991	54.747	54.678	61.116	58.535	131.703	97.692
	Y79AA1000989	160.869	133.278	169.716	48.057	196.947	67.040	105. 199	90.492
40	Y79AA1000991	172.775	159.227	83.980	68.958	59.956	152.374	108.299	84.387
	Y79AA1001013	199, 195	153.480	107.292	61.287	92.604	113.848	154. 343	119.100
	Y79AA1001014	68.728	72.126	41.236	31.089	17.667	51, 104	41, 121	35. 352
	Y79AA1001019	66.003	34.676	36. 574	22.751	21.527	33.525	40.467	35.925
	Y79AA1001020	58.188	33.720	31.511	41.189	21.352	33.976	46.407	37.451
	Y79AA1001023	75.610	41.776	31.044	17. 988	30.650	42.942	60.331	30, 561
45	Y79AA1001030	103. 273	36.017	33.752	31.467	19.917	43.990	63. 269	32.983
	Y79AA1001035	0.000	0.000	28.444	28.051	16.127	41.569	62.544	47.884
	Y79AA1001041	77.214	55, 578	30.400		26.174	46.066	33.311	28.914
	Y79AA1001043	62, 920	86.930	40. 257	39.379	42.525	44, 192	65.573	44.307
	Y79AA1001048	69, 373	57, 191	47.559	29.744	25. 491	59.541	61.196	33.290
	Y79AA1001056	28. 105	21.448	25.068			27. 941	27.218	31.106
50	Y79AA1001061	77.662	63, 993	57.624			42.698	30.186	47. 071 19. 189
-	Y79AA1001062	23.211	15. 295	22.974			12.268 47.635	15. 522 42. 292	
	Y79AA1001068	89.610	80.709 77.800	62.102					
	Y79AA1001073	167.563	91.034		52. 366				65. 133
	Y79AA1001077	128.286	31.034	. 02.331	1 32. 300	1 11.143	1. 1.5. 552	1 .55.0.7	

Table 166

								O 1 1 I	26.651
	Y79AA1001078	23.435	19. 289	16.494	16.707	8. 916	16.759	28,013	25.651
	Y79AA1001081	80.143	68.142	45.763	36.383	26.159	35.757	38.026	35.885
			242.117		24.084	134,444	174.586	238.334	149.593
	Y79AA1001088	317.039				55, 190	98. 837	117.534	77.578
5	Y79AA1001089	198.139	98.655	80.498	49.545			32.352	35.391
	Y79AA1001090	80.451	60.910	39.633	42.380	36.692	37. 452		
	Y79AA1001105	242.673	66.561	63.208	31.037	76.586	75, 243	109.216	60.833
		79.091	23.396	18.843	28.396	19.935	55.429	96.508	34. 254
	Y79AA1001142		201.081		08.956	107.663	126.922	147.749	112, 199
	Y79AA1001145	227.540			13.823	7.016	14, 608	10.831	11.076
	Y79AA1001162	32.474	21. 215	17. 402				27.977	21.861
10	Y79AA1001167	81.840	38. 276	27. 439	20.713	20. 465	39, 401		
. •	Y79AA1001176	37.234	30. 174	29.821	28.145	17, 772	23.084	23.905	31.875
	Y79AA1001177	157.278	72.492	47.515	31,006	45, 407	62. 162	74.915	44.631
			77.734	66. 981	49.326	60.911	108, 763	101.419	45.761
	Y79AA1001179	155. 289			18.392	18. 203	25, 381	22.095	14. 576
	Y79AA1001185	42.293	30. 499	20.818			55. 945	44, 441	55. 417
	Y79AA1001201	70. 267	62.245	55. 927	64.637	42.307			
15	Y79AA1001205	76.691	73.411	29.446	25.089	10.867	25. 196	31,540	23.771
13	Y79AA1001211	69.077	77. 295	43.109	54,773	26. 171	19.436	23.382	36.508
			40.760	30.464	21.472	22.536	28.939	31.790	27. 996
	Y79AA1001212	60.509				48.098	86.493	78.651	128.332
	Y79AA1001216	107.414	112.384,	51.845	90. 341			95.311	72.216
	Y79AA1001228	191.014	98.191	77.471	55.138	68.036	114. 392		
	Y79AA1001233	165. 200	46.959	55.748	19, 356	50.639	93. 326	77.766	29.974
	Y79AA1001235	75.419	41.716	32.067	19. 238	31.896	34.830	44, 490	38.856
20	Y79AA1001239	348. 195	155. 335	206. 398	93. 364	264.580	150. 282	141. 282 -	138.685
			55. 824	32.015	19. 335	24. 480	129.654	123.682	27.590
	Y79AA1001240	97.619				23, 430	42.093	44. 389	40,863
	Y79AA1001255	60, 196	39.594	29.713	32.087		20, 450	18.219	19.822
	Y79AA1001264	23.500	30.229	13.518	13.380	8, 385			
	Y79AA1001272	172.136	148.159	89.874	101.905	67.677	109. 162	89.962	89.461
	Y79AA1001281	23. 625	18.360	9, 518	9.700	6. 169	17.324	15.120	11.543
25	Y79AA1001299	257. 530	138.510	106.642	92, 167	96, 141	155.017	156.902	114.884
			18.932	11.140	5, 860	16.123	10, 337	9, 558	9.283
	Y79AA1001312	28. 599				80.506	137.595	117. 523	59, 456
	Y79AA1001319	233. 396	111.817	90. 283	51.100			31.769	19.583
	Y79AA1001323	46. 240	62.299	28.364	20.915	20.142	36.013		
	Y79AA1001328	166. 188	85. 958	71.107	51.952	47.867	98.151	92.634	63.952
	Y79AA1001343	5293. 557	1957.671	5529.524	508.017	5447.748	5598. 173	4563.395	1662.056
22		23.608	13.189	12.127	7,610	6. 082	11,346	6.319	6.967
30	Y79AA1001351				44.078	18.806	18.623	17.892	AAA
									1 57.833
	Y79AA1001364	23.462	34.748	26. 228					57.833
	Y79AA1001367	74.110	39.168	25. 534	16.038	21, 213	33.215	35, 782	29.409
	Y79AA1001367				16.038 6.267	21, 213 19, 195	33. 215 15. 742	35, 782 34, 303	29.409 21.015
	Y79AA1001367 Y79AA1001384	74.110 44.135	39.168 26.692	25. 534	16.038	21, 213	33.215 15.742 38.414	35, 782 34, 303 60, 920	29. 409 21. 015 32, 481
	Y79AA1001367 Y79AA1001384 Y79AA1001391	74.110 44.135 88.486	39.168 26.692 45.427	25. 534 19. 494 33. 937	16. 038 6. 267 20. 520	21, 213 19, 195 35, 938	33. 215 15. 742 38. 414 26. 264	35, 782 34, 303 60, 920 37, 409	79. 409 21. 015 32. 481 30. 457
	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001394	74.110 44.135 88.486 73.046	39.168 26.692 45.427 48.196	25. 534 19. 494 33. 937 27. 660	16.038 6.267 20.520 20.614	21, 213 19, 195 35, 938 16, 092	33. 215 15. 742 38. 414 26. 264	35, 782 34, 303 60, 920 37, 409	29. 409 21. 015 32, 481
35	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001394 Y79AA1001402	74.110 44.135 88.486 73.046 277.943	39. 168 26. 692 45. 427 48. 196 171. 103	25. 534 19. 494 33. 937 27. 660 185. 389	16.038 6.267 20.520 20.614 101.994	21, 213 19, 195 35, 938 16, 092 137, 576	33. 215 15. 742 38. 414 26. 264 164. 575	35, 782 34, 303 60, 920 37, 409 126, 561	29.409 21.015 32.481 30.457 96.457
35	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001394 Y79AA1001402 Y79AA1001410	74.110 44.135 88.486 73.046 277.943 37.405	39.168 26.692 45.427 48.196 171.103 47.535	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875	16.038 6.267 20.520 20.614 101.994 21.151	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753	33.215 15.742 38.414 26.264 164.575 29.322	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883
35	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001402 Y79AA1001410 Y79AA1001414	74.110 44.135 88.486 73.046 277.943 37.405 40.424	39.168 26.692 45.427 48.196 171.103 47.535 18.548	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585	16.038 6.267 20.520 20.614 101.994 21.151 8.705	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753 9, 528	33.215 15.742 38.414 26.264 164.575 29.322 27.024	35, 782 34, 303 60, 920 37, 409 126, 561 20, 709 20, 661	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409
35	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001394 Y79AA1001402 Y79AA1001410	74.110 44.135 88.486 73.046 277.943 37.405	39.168 26.692 45.427 48.196 171.103 47.535	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982	16.038 6.267 20.520 20.614 101.994 21.151 8.705 17.958	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753 9, 528 30, 855	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863	35, 782 34, 303 60, 920 37, 409 126, 561 20, 709 20, 661 82, 932	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756
35	Y79A1001367 Y79A1001384 Y79A1001391 Y79A1001394 Y79A1001402 Y79A1001410 Y79A1001414 Y79A1001426	74.110 44.135 88.486 73.046 277.943 37.405 40.424	39.168 26.692 45.427 48.196 171.103 47.535 18.548	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585	16.038 6.267 20.520 20.614 101.994 21.151 8.705	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753 9, 528 30, 855 42, 573	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818	35, 782 34, 303 60, 920 37, 409 126, 561 20, 709 20, 661 82, 932 79, 641	79, 409 21, 015 32, 481 30, 457 96, 457 19, 883 18, 409 35, 756 62, 907
35	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001402 Y79AA1001410 Y79AA100146 Y79AA1001427	74.110 44.135 88.485 73.046 277.943 37.405 40.424 128.039	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982	16.038 6.267 20.520 20.614 101.994 21.151 8.705 17.958	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753 9, 528 30, 855	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825	35, 782 34, 303 60, 920 37, 409 126, 561 20, 709 20, 661 82, 932 79, 641 47, 406	79, 409 21, 015 32, 481 30, 457 96, 457 19, 883 18, 409 35, 756 62, 907 24, 141
35	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001410 Y79AA1001426 Y79AA1001427 Y79AA1001427 Y79AA1001430	74.110 44.135 88.486 73.046 277.943 37.405 40.424 128.039 102.517 88.291	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775	16.038 6.267 20.520 20.614 101.994 21.151 8.705 17.958 26.901 15.144	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753 9, 528 30, 855 42, 573 26, 578	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818	35, 782 34, 303 60, 920 37, 409 126, 561 20, 709 20, 661 82, 932 79, 641	79, 409 21, 015 32, 481 30, 457 96, 457 19, 883 18, 409 35, 756 62, 907
	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001414 Y79AA1001426 Y79AA1001426 Y79AA1001430 Y79AA1001430	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643	16.038 6.267 20.520 20.614 101.994 21.151 8.705 17.958 26.901 15.144 11.993	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792	79, 409 21, 015 32, 481 30, 457 96, 457 19, 883 18, 409 35, 756 62, 907 24, 141
35	Y79A1001367 Y79A1001384 Y79A1001394 Y79A1001394 Y79A1001410 Y79A1001410 Y79A1001414 Y79A1001427 Y79A1001427 Y79A1001433 Y79A1001485	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416	16. 038 6. 267 20. 520 20. 614 101. 994 21. 151 8. 705 17. 958 26. 901 15. 144 11. 993 6. 180	21, 213 19, 195 35, 938 16, 092 137, 576 18, 753 9, 528 30, 855 42, 573 26, 578 13, 637 7, 239	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923 8.477	35, 782 34, 303 60, 920 37, 409 126, 561 20, 709 20, 661 82, 932 79, 641 47, 406 33, 792 11, 343	79, 409 21, 015 32, 481 30, 457 96, 457 19, 883 18, 409 35, 756 62, 907 24, 141 21, 060 9, 667
	Y79AA1001367 Y79AA1001384 Y79AA1001394 Y79AA1001394 Y79AA1001410 Y79AA1001410 Y79AA1001427 Y79AA1001427 Y79AA1001439 Y79AA1001439 Y79AA1001485 Y79AA1001485	74.110 44.135 88.486 73.046 277.943 37.405 40.424 128.039 102.517 88.291 22.600 12.457 3.325	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808	16. 038 6. 267 20. 520 20. 614 101. 994 21. 151 8. 705 17. 958 26. 901 15. 144 11. 993 6. 180 0. 913	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923 8.477 2.370	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535
	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001402 Y79AA1001410 Y79AA1001414 Y79AA1001427 Y79AA1001430 Y79AA1001430 Y79AA1001430 Y79AA1001430 Y79AA1001433	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 42.870	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800	16. 038 6. 267 20. 520 20. 514 101. 994 21. 151 8. 705 17. 958 26. 901 15. 144 11. 993 6. 180 0. 913 22. 668	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923 8.477 2.370	35. 782 34. 303 60. 920 37. 409 126. 561 20. 769 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671
	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001402 Y79AA1001410 Y79AA1001414 Y79AA1001427 Y79AA1001430 Y79AA1001430 Y79AA1001430 Y79AA1001430 Y79AA1001433	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808	16. 038 6. 267 20. 520 20. 614 101. 994 21. 151 8. 705 17. 958 26. 901 15. 144 11. 993 6. 180 0. 913	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 755 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232
	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001416 Y79AA1001426 Y79AA1001427 Y79AA1001430 Y79AA1001439 Y79AA1001439 Y79AA1001431 Y79AA1001431 Y79AA1001511 Y79AA1001511	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 337 131. 638	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923 8.477 2.370 30.907 38.384 26.821	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 5375 41. 671 16. 232 15. 958
	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001414 Y79AA1001426 Y79AA1001439 Y79AA1001439 Y79AA1001439 Y79AA1001493 Y79AA1001523 Y79AA1001523	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 42.870 41.082	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094
40	Y79AA1001367 Y79AA1001384 Y79AA1001394 Y79AA1001400 Y79AA1001410 Y79AA1001416 Y79AA1001427 Y79AA1001427 Y79AA1001439 Y79AA1001485 Y79AA1001511 Y79AA1001513 Y79AA1001533	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 42.870 41.082 17.602 81.487	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 7775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 11.993 6.180 0.913 22.668 24.376 9.778	21. 213 19. 195 35. 938 16. 052 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 33. 229 52. 833	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923 8.477 2.370 30.907 38.384 26.821 44.930	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 5375 41. 671 16. 232 15. 958
	779A1001367 779A1001384 779A1001391 779A1001391 779A1001402 779A1001410 779A1001410 779A1001427 779A1001427 779A1001427 779A1001439 779A1001511 779A1001523 779A1001523 779A1001523 779A1001530 779A1001530	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 71. 806	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082 17. 602 81. 487 80. 795	25. 534 19. 454 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 31. 800 28. 617 37. 936 57. 603 31. 639	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296 34.117	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 30. 907 38. 384 26. 821 44. 930 30. 573	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551
40	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001410 Y79AA1001414 Y79AA1001430 Y79AA1001430 Y79AA1001430 Y79AA1001433 Y79AA1001433 Y79AA1001513 Y79AA1001523 Y79AA1001532 Y79AA1001532 Y79AA1001532	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 71. 806 21. 702	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 42.870 41.082 17.602 23.664	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 31. 639 13. 568	16.038 6.267 20.520 20.514 100.994 21.151 3.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 7. 239 30. 480 38. 748 39. 229 52. 833 34. 465	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 19. 043	35. 782 34. 303 60. 920 37. 409 126. 561 20. 769 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890
40	779A1001367 779A1001384 779A1001391 779A1001391 779A1001402 779A1001410 779A1001410 779A1001427 779A1001427 779A1001427 779A1001439 779A1001511 779A1001523 779A1001523 779A1001523 779A1001530 779A1001530	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 71. 806 21. 702	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082 17. 602 81. 487 80. 795 23. 664 125. 939	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 13. 558 91. 450	16.038 6.267 20.520 20.614 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296 33.413 78.443	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 19. 043	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890
40	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001410 Y79AA1001414 Y79AA1001430 Y79AA1001430 Y79AA1001430 Y79AA1001433 Y79AA1001433 Y79AA1001513 Y79AA1001523 Y79AA1001532 Y79AA1001532 Y79AA1001532	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 21. 702	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082 17. 602 81. 487 80. 795 23. 664 125. 939	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 11. 558 91. 450	16.038 6.267 20.520 20.614 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296 33.413 78.443	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 102. 345 64. 477	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245
40	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001391 Y79AA1001410 Y79AA1001416 Y79AA1001426 Y79AA1001439 Y79AA1001439 Y79AA1001439 Y79AA1001530 Y79AA1001530 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001534	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 71. 806 21. 702 160. 862	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 41.082 17.602 81.487 80.795 23.664 125.939 77.112	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 37. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 31. 639 91. 450 44. 627	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 11.993 6.180 0.913 22.686 24.376 9.778 49.296 34.117 13.443 78.443 26.543	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 19. 644	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245
40	779AA1001367 779AA1001384 779AA1001394 779AA1001402 779AA1001410 779AA1001427 779AA1001427 779AA1001427 779AA1001427 779AA1001430 779AA1001511 779AA1001523 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 71. 806 21. 702 154. 131	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082 81. 487 80. 795 23. 664 125. 939 77. 112	25. 534 19. 454 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 57. 603 31. 639 13. 558 91. 450 44. 627 21. 323	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296 34.117 13.443 78.443 78.443 25.543	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269 21. 126	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 44. 930 30. 573 19. 643 102. 345 64. 477 20. 650	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 77. 908	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245
40	Y79AA1001367 Y79AA1001384 Y79AA1001391 Y79AA1001391 Y79AA1001402 Y79AA1001410 Y79AA1001427 Y79AA1001427 Y79AA1001427 Y79AA1001439 Y79AA1001523 Y79AA1001523 Y79AA1001523 Y79AA1001523 Y79AA1001530 Y79AA1001541 Y79AA1001541 Y79AA1001541 Y79AA1001541 Y79AA1001541 Y79AA1001542 Y79AA1001541	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 71. 806 21. 702 160. 862 154. 131 19. 278 29. 260	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 42.870 41.082 17.602 81.487 81.487 82.3664 125.939 77.112	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 31. 639 13. 558 91. 450 44. 627 21. 323 4. 472	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296 34.117 13.443 78.443 25.543 15.645 2.139	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 21. 126 6. 817	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 19. 043 102. 345 64. 477 20. 650 9. 566	35. 782 34. 303 60. 920 37. 409 126. 561 20. 769 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 72. 908 81. 174 8. 517	29. 409 21. 015 32. 481 30. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245 27. 432 4. 043
40	Y79AA1001367 Y79AA1001384 Y79AA1001394 Y79AA1001402 Y79AA1001410 Y79AA1001410 Y79AA1001430 Y79AA1001430 Y79AA1001439 Y79AA1001439 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 21. 702 160. 862 154. 131 2. 19. 278	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082 17. 602 81. 487 77. 112 39. 676 125. 939 77. 112 39. 676 9. 273	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 31. 639 13. 558 91. 450 44. 627 21. 323 4. 472 5. 154	16.038 6.267 20.520 20.514 101.994 21.151 3.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.296 34.117 13.443 78.443 26.543 15.462 2.139 4.611	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269 21. 126 6. 817 8. 363	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 19. 043 102. 345 64. 477 9. 566 9. 566	35. 782 34. 303 60. 920 37. 409 126. 561 20. 769 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 72. 908 8. 517 8. 575	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245 40. 433 4. 043 6. 599
40 45	Y79AA1001367 Y79AA1001384 Y79AA1001394 Y79AA1001402 Y79AA1001410 Y79AA1001416 Y79AA1001426 Y79AA1001430 Y79AA1001430 Y79AA1001433 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001546 Y79AA1001555 Y79AA1001555 Y79AA1001555 Y79AA1001555 Y79AA1001555	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 21. 702 160. 862 154. 131 19. 278 29. 266 10. 832	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 41.082 17.602 81.487 42.870 41.082 17.602 81.487 125.939 77.112 39.676 1.846 9.273 61.837	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 31. 639 13. 558 91. 450 44. 627 21. 323 44. 472 5. 154 49. 013	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.686 24.376 9.778 49.296 34.117 13.443 78.443 26.543 15.462 2.139 4.611 45.123	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269 21. 126 6. 817 8. 363 38. 746	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 30. 907 38. 384 26. 821 44. 930 102. 345 64. 477 20. 650 3. 849 42. 497	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 72. 908 11. 744 8. 517 8. 515	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 755 62. 907 24. 141 21. 060 9. 667 15. 355 41. 671 16. 232 15. 958 47. 094 47. 094 47. 551 17. 890 92. 148 37. 245 27. 432 6. 599 62. 013
40 45	Y79AA1001367 Y79AA1001384 Y79AA1001394 Y79AA1001402 Y79AA1001410 Y79AA1001410 Y79AA1001430 Y79AA1001430 Y79AA1001439 Y79AA1001439 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 21. 702 160. 862 154. 131 19. 278 29. 266 10. 832	39.168 26.692 45.427 48.196 171.103 47.535 18.548 45.365 75.088 44.524 31.240 15.003 3.087 41.082 17.602 81.487 42.870 41.082 17.602 81.487 125.939 77.112 39.676 1.846 9.273 61.837	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 31. 639 13. 558 91. 450 44. 627 21. 323 44. 472 5. 154 49. 013	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.686 24.376 9.778 49.296 34.117 13.443 78.443 78.443 15.462 2.139 45.123 20.106	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269 21. 126 6. 817 8. 363 38. 746 22. 571	33.215 15.742 38.414 26.264 164.575 29.322 27.024 79.863 49.818 35.825 8.923 8.477 2.370 30.907 38.384 26.821 44.930 30.573 19.643 102.345 64.477 20.650 9.566 3.848 42.497	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 82. 932 79. 641 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 72. 908 11. 744 8. 517 8. 753 50. 134	79. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 35. 756 62. 907 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245 27. 432 4. 043 37. 245 27. 432 4. 043 34. 587
40 45	779AA1001367 779AA1001384 779AA1001384 779AA1001394 779AA1001402 779AA1001410 779AA1001427 779AA1001427 779AA1001427 779AA1001511 779AA1001523 779AA1001523 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532 779AA1001532	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 71. 806 21. 702 160. 862 154. 131 19. 278 29. 260 10. 832 95. 166	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 41. 082 17. 602 81. 487 80. 795 23. 664 125. 939 77. 112 39. 676 9. 273 61. 837 50. 427	25. 534 19. 494 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 17. 775 12. 643 6. 416 0. 808 31. 800 28. 617 37. 936 57. 603 31. 639 91. 450 44. 627 21. 323 4. 472 49. 013 16. 817	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 11.993 6.180 0.913 22.686 24.376 9.778 49.296 34.117 13.443 78.443 78.443 26.543 15.462 2.139 4.611	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269 21. 126 6. 817 8. 363 38. 746 22. 571	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 19. 043 102. 345 64. 477 20. 650 9. 565 3. 889 42. 497 18. 261	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 72. 908 11. 744 8. 517 8. 753 50. 134 3. 5915	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245 27. 432 4. 043 6. 599 62. 013 34. 587 148. 708
40 45	Y79AA1001367 Y79AA1001384 Y79AA1001394 Y79AA1001402 Y79AA1001410 Y79AA1001416 Y79AA1001426 Y79AA1001430 Y79AA1001430 Y79AA1001433 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001532 Y79AA1001546 Y79AA1001555 Y79AA1001555 Y79AA1001555 Y79AA1001555 Y79AA1001555	74. 110 44. 135 88. 486 73. 046 277. 943 37. 405 40. 424 128. 039 102. 517 88. 291 22. 600 12. 457 3. 325 34. 387 131. 638 64. 263 84. 756 71. 806 21. 702 160. 862 154. 131 19. 278 29. 260 10. 812 195. 166	39. 168 26. 692 45. 427 48. 196 171. 103 47. 535 18. 548 45. 365 75. 088 44. 524 31. 240 15. 003 3. 087 42. 870 41. 082 81. 487 80. 795 23. 664 125. 939 77. 112 39. 676 1. 846 9. 273 50. 427	25. 534 19. 454 33. 937 27. 660 185. 389 22. 875 20. 585 44. 982 38. 728 17. 775 12. 643 31. 800 28. 617 37. 936 57. 603 31. 639 13. 558 91. 450 44. 627 21. 323 4. 472 5. 154	16.038 6.267 20.520 20.514 101.994 21.151 8.705 17.958 26.901 15.144 11.993 6.180 0.913 22.668 24.376 9.778 49.295 34.117 13.443 78.443 26.543 15.462 2.139 4.611 45.123 70.106 89.900	21. 213 19. 195 35. 938 16. 092 137. 576 18. 753 9. 528 30. 855 42. 573 26. 578 13. 637 7. 239 0. 895 30. 480 38. 748 39. 229 52. 833 34. 465 17. 622 96. 657 33. 269 21. 126 6. 817 8. 363 38. 748 39. 229 50. 835 30. 855 40. 578 40. 578	33. 215 15. 742 38. 414 26. 264 164. 575 29. 322 27. 024 79. 863 49. 818 35. 825 8. 923 8. 477 2. 370 30. 907 38. 384 26. 821 44. 930 30. 573 19. 043 102. 345 64. 477 20. 650 9. 566 3. 849 42. 497 18. 261 86. 111	35. 782 34. 303 60. 920 37. 409 126. 561 20. 709 20. 661 47. 406 33. 792 11. 343 3. 288 27. 949 55. 678 36. 155 58. 976 56. 137 24. 725 81. 132 72. 908 11. 744 8. 517 8. 753 50. 134 3. 5915	29. 409 21. 015 32. 481 30. 457 96. 457 19. 883 18. 409 24. 141 21. 060 9. 667 1. 535 41. 671 16. 232 15. 958 47. 094 27. 551 17. 890 92. 148 37. 245 27. 432 4. 043 6. 599 62. 013 34. 587 148. 708

Table 167

	CONTRACTOR I	13.646	12.156	6.553	6.307	8.775	5. 570 T	14.006	7.900
	Y79AA100163D								
	Y79AA1001647	43.380	30.209	38. 642	18.607	50.749	13.890	27.635	18.826
	Y79AA1001664	50.619	62.037	23. 222	35.535	31.203	25.999	33. 586	35.257
5	Y79AA1001665	78.815	50.214	28, 199	20.230	28. 531	39. 239	43.686	24.873
	Y79AA1001679	182.502	59.845	49, 481	32, 964	56.908	102, 379	88.857	48.625
		48.740	44.701	21. 354	18,732	23.271	23.639	35.010	24.377
	Y79AA1001692					14.623	7.312	6.730	10.898
	Y79AA1001696	6,780	14. 124	10.007	8.631				
	Y79AA1001705	84.869	54. 294	35.569	21. 435	39. 991	44.064	55.777	37. 387
	Y79AA1001711	62.806	75.073	36.984	31. 331	32.851	38. 989	52.758	53.508
10		21.280	34. 089	12.028	10.889	15.395	11.748	24. 174	10.438
10	Y79AA1001717		51.690	17.623	19.362	20.441	20.351	30. 934	27.264
	Y79AA1001719	43.417				54.868	25.062	43. 179	39.080
	Y79AA1001727	73.341	78.139	42.958	28.661		113.999	151, 389	151, 480
	Y79AA1001750	294.250	240.534		113.859	147. 591			
	Y79AA1001760	186.817	180.985	65.822	68.036	91.745	92. 228	144. 742	103.455
	Y79AA1001777	125.250	87.579	38. 902	31.498	34.787	51. 175	61.299	47. 365
	Y79AA1001781	0.000	0.000	0.000	3.210	1.473	0,000	0.000	0.000
15			61.186	49.706	30.708	31.661	57.179	72.608	156. 355
	Y79AA1001787	114.565				48, 463	74. 105	167.655	89.090
	Y79AA1001793	185.933	88.770	84.898	59.826				
	Y79AA1001795	17.050	21.582,	20. 234	15.314	13.998	18.815	13.699	16.861
	Y79AA1001799	86.419	58. 330	51.694	35. 379	30,638	47, 701	71.948	46.535
	Y79AA1001800	511.812	97. 958	354.971	49. 190	235. 401	413. 238	490. 565	73.961
	Y79AA1001801	67.645	45.550	11.683	27.943	20.138	30. 425	59. 231	35. 253
20		57. 387	56.011	59. 286	17.563	64.207	39. 997	39. 216	36.149
	Y79AA1001803			125. 538	76. 023	66. 375	74, 500	58. 882	68.354
	Y79AA1001805	146.263	170.531				46. 384	86. 296	39. 428
	Y79AA1001807	112.057	63.466	54.660	28.588	34. 253			
	Y79AA1001827	70.024	30. 424	44.198	33.684	23.703	26. 135	51.859	17.610
	Y79AA1001846	25.975	42.461	56.527	62. 241	32.960	50. 520	20.001	36.949
	Y79AA1001848	35.746	22.982	23, 160	9.894	16.543	12.462	26.092	21.091
25	Y79AA1001853	281.071	150.082	159.752	107,770	164. 169	199.036	174. 168	111.109
23			108.799	96. 407	63.758	66.145	100.694	163.628	77.595
	Y79AA1001863	190.420			28. 167	24. 388	28.450	20.721	74. 899
	Y79AA1001866	24, 530	46.991	37.466				1.506	1. 497
	Y79AA1001874	1.221	5. 487	0.848	1.231	0, 291	0. 598		
	Y79AA1001875	63.952	58.462	47. 435	36.846	24. 598	39.313	45. 106	40. 636
	Y79AA1001907	124.410	250.090	50.333	92.943	49.772	74. 402	107.811	194. 562
	Y79AA1001908	12.574	13.547	9.612	6.931	5. 169	7.911	11.534	9.867
30	Y79AA1001923	33.869	14. 234	14, 248	5.718	8, 352	12.798	25. 326	7.829
				44.024	41.115	46, 490	154. 335	107.236	39. 239
	Y79AA1001927	186.717	76.975			15.694	32. 271	26. 423	29.042
	Y79AA1001930	33.259	33.470	18.855	24. 182			10. 994	25.644
	Y79AA1001932	27.741	23. 277	12.768	9, 914	14.699	8. 522		
	Y79AA1001933	34.948	36.160	27.478	18, 608	18. 230	17. 284	30. 314	30.361
	Y79AA1001942	28.803	28. 251	22.497	11.034	11.547	51.771	43.263	5. 042
35	Y79AA1001963	68. 323	43.878	42.080	36, 240	33.736	26.445	62.945	56.785
	Y79AA1001968	55. 189	120. 287	31, 107	72.431	32.780	37. 209	52. 124	87.863
			44, 245	40.209	17, 481	29. 219	49.886	55. 561	26, 162
	Y79AA1001983	91.447				28.700	41.938	31, 511	26.090
	Y79AA1002000	78.569	42.344	37.253	28.054	25. 700	62.884	62.767	36.527
	Y79AA1002004	135.629	61.297	65.308	50.333	46.897			
	Y79AA1002008	151. 334	65.665	44.780	33, 954	37. 173	46, 166	78. 471	49. 925
40	Y79AA1002012	140. 300	132.533	88. 285	105.977	78.145	59.701	57. 183	104. 179
	Y79AA1002017	38. 327	28.859	17.564	7. 197	8. 297	22.866	39. 108	18. 179
	Y79AA1002022	197.012	109.640	111.812	71.115	68.794	122.840	108. 973	82.258
	Y79AA1002027	7.861	6.807	5.719	3, 405	5. 503	3.619	4. 936	6.690
	Y79AA1002050	52.645	57.007	34. 182	49, 101	29.451	26.571	25, 198	32.168
					46.043	84. 452	92, 949	131.501	99.159
	Y79AA1002058	162.814	86.786	63, 856			27.031	33, 624	43.906
45	Y79AA1002060	74.517	43. 157	38.911	33.852	57.622			
	Y79AA1002082	163.546	122. 645	81.975	88.856	64. 753	92. 455	75. 321	73. 162
	Y79AA1002065	72.537	83.880	24.771	44. 298	22.044	30.756	35. 287	75. 559
	Y79AA1002067	18.914	16.681	8.561	6.098	4. 972	7.856	26, 231	10.844
		153.130	40.848	44, 030	9.535	26. 886	75.515	76.585	29.038
	Y79AA1002069			64, 787	58, 175	89. 400	172.062	64. 355	62.998
	Y79AA1002070	255.333	63.953			107.873	170.520	153.058	388. 635
50	Y79AX1002074	168.399	367.145	81.099					
50	Y79AA1002076	36.931	26.480	13.779			20.354	19. 122	14.650
	Y79AA1002083	100.267	39. 527	25. 359			42.095	30.686	16.092
	Y79AA1002084	31.602	37. 320	24.313	14.210	13, 535	21.829	27.098	16.412
	Y79AA1002086	43.060		21.971			19.023	17.056	19.318
		1 .0. 230							

Table 168

				1 20	16 109			_	
	Y79AA1002087	13.030	15. 226	11.425	22. 378	3.745	12.088	10.009	24. 908
	Y79AA1002089	40.323	26. 458	12.982	15.098	16.218	17.576	3.691	24.665
	Y79AA1002093	46, 120	27.022	18, 769	15, 919	7.245	24.041	28. 202	24.994
5	Y79AA1002101	43.837	30.418	18.385	11.894	11.521	24.278	23. 182	15.994
	Y79AA1002103	43. 141	24. 675	23. 246	31.726	17. 340	31, 371	32. 322	48. 954
		20.756	22, 498	17.048	10.575	15. 180	11.669	14.011	12.945
	Y79AA1002115	27. 091	49. 228	19. 524	15. 594	14.827	12, 987	19, 044	18. 216
	Y79AA1002121		64, 875	48. 646	23, 137	27. 474	32.479	34. 123	51.389
	Y79AA1002125	48.808	25. 472	14. 117	14. 375	7. 485	13. 555	11.317	12.960
10	Y79AA1002129	20.607	22.411	18, 720	14, 115	7.829	20. 162	15. 429	12.884
	Y79AA1002131	46. 336	11, 713	5.758	6.335	6. 389	11, 186	2.854	4. 539
	Y79AA1002139	17. 296	47.677	66.378	20.967	59.407	32. 426	31. 597	21. 322
	Y79AA1002144	45. 269 301. 285	121.825	100.055	57.536	81.697	176. 423	154, 681	88. 082
	Y79AA1002177		99. 397	37.780	10.625	40.969	35, 101	65.850	66. 184
	Y79AA1002183	78.011	69.118	26.355	26. 998	31, 172	30.882	39, 528	28. 104
15	Y79AA1002202	\$7.948	53,775	45.674	14.730	26, 902	42. 785	47. 433	32.007
	Y79AA1002204	108. 226	20,653	11, 579	11, 189	8, 007	20. 198	14.716	14.423
	Y79AA1002206	23.882	19.145	14. 805	15. 985	9, 456	19.745	11.177	17.666
	Y79AA1002208	17, 539		11. 592	3.770	5. 884	7.681	9. 212	6.769
	Y79AA1002209	12.404	10.671,	11. 197	4. 453	8. 279	31.518	24. 637	13. 120
	Y79AA1002210	36.693	21.704	23. 317	18.415	22, 277	33. 188	47.655	53.021
20	Y79AA1002211	60.744 88.865	40.012 56.933	24. 906	28.654	40, 420	32. 547	31. 240	41.587
	Y79AA1002213		74, 421	32.504	25. 568	33. 392	47.741	30, 830	34, 812
	Y79AA1002215	57.323 7.686	27.673	7. 325	5. 327	8. 309	5. 571	9. 728	9. 037
	Y79AA1002220 Y79AA1002226	33.811	70. 351	53.822	44. 542	43, 103	43.566	31.798	56,096
		133.812	49.906	27.621	14.021	32.478	73.121	60. 968	21.211
	Y79AA1002229 Y79AA1002234	53.796	27. 231	31.097	16.258	22. 352	39. 228	41.686	31.562
25	Y79AA1002235	9.109	6.947	3. 938	3, 201	5.077	8.688	8.099	8. 031
	Y79AA1002246	45.749	34.031	22,771	19, 593	19, 245	14.798	40.274	41.271
	Y79AA1002258	75, 546	58.416	30.618	24.590	30. 971	35.864	47.893	50.632
	Y79AA1002279	67.007	468.054	23.705	27. 332	22.243	72.113	23.817	64.255
	Y79AA1002292	107.375	48.724	45. 677	27.662	41. 581	54.031	48.041	36.807
	Y79AA1002298	16.948	16.878	8.834	7. 151	8,601	7.054	11.871	9. 334
30	Y79AA1002307	29.343	26.868	16.693	17. 533	20, 451	13. 735	13.467	11.704
	Y79AA1002309	38.982	33.605	15. 626	14. 434	15. 282	17.723	25. 386	17. 397
	Y79AA1002311	31.668	30.875	21.323	22, 152	19. 332	10.916	32.170	15. 265
	Y79AA1002334	49. 431	32.284	18. 242	13.025	24.412	19.450	30.870	24. 306
	Y79AA1002351	41, 486	18,773	27.420	13. 424	23.100	22. 549	45. 251	26.383
35	Y79AA1002355	10.396	23.208	37.472	13.874	42.683	14.865	12.092	15. 185
33	Y79AA1002361	88.085	78.594	36.358	37.149	35.845	41.778	36.660	25. 294
	Y79AA1002365	17.588	21.447	10. 949	7.231	11.431	16.111	15, 168	14. 782
	Y79AA1002373	50.748	39.981	17.086	11.669	21, 120	12.396	22.757	15. 438
	Y79AA1002376	6643.977		4553. 953	585. 102	6666.479	5319.310	6496.197	1220.015
	Y79AA1002378	77.584	97.591	29. 238	27. 161	35.356	35. 168	42.325	47. 261
40	Y79AA1002381	141.196	111.531	39. 904	43.874	44.814	57.151	75.416	73. 250
	Y79AA1002388	165.548	86.006	56. 942	27. 181	60.647	43.749	87.173 42.014	61.931
	Y79AA1002399	47. 127	38. 224	20.037	14.800	14. 138	25. 545 15. 977		15.674
	Y79AA1002407	14.750	20.995	15. 394	14.318	11.321	85. 902	15.721 42.923	62.238
	Y79AA1002413	55.733	94.994	61.674	38. 953	26.005	17. 322	34. 129	20.886
	Y79AA1002416	26.021	26.133	18.893	17.489		12.675	14.708	62. 243
45	Y79AA1002429	29. 180	51.475	14.818	24. 101	19, 762	32. 188		19, 906
	Y79AA1002431	36.374	37.521	29.072			44, 782	56.569	48.003
	Y79AA1002433	73.392	56.725	40.689	45.773	29.753 84.172	187. 480	65.701	66.873
	Y79AA1002445	206.082	130.492	119. 284	81.825	30.726	58.954	73.797	51.203
	Y79AA1002461	136. 322	87.178	56. 327	41.540	27.818	62.743	46. 169	48.544
	Y79AA1002466	58.450	66.910	32.039	63.994	17.619	16.828	33.234	22.949
50	Y79AA1002471	22. 153	38.198	21.750	19.098	43.775	40. 296	44.510	52.633
	Y79AA1002472	60. 980	65.699	60. 101	81.738	17.029	18.872	29.379	12.444
	Y79AA1002474	35. 222	8. 126	16.456	10.777	82. 291	51.719	43. 236	91.558
	Y79AA1002482	72.994	104.184	83. 915	153, 120	9.046	9. 098	17. 186	12.270
	Y79AA1002487	22.033	18.529 63.572	10.754	10, 800	29.453	73.670	63. 467	29.681
	Y79AA1002490	105.735	80. 901	47. 379		35.093	37.512		19. 166
55	779AA1002493 ZRV6C1006278	72.446 37.372	7. 298	8. 151	3.848	4, 644		6.612	2.417
	THIST INDELIG	1 31.312	1. 230	1 3. 131	1 3.040	7.074	<del></del>		

# Table 169

Expression of each cDNA in human pulmonary arterial endothelial cells cultured in a medium containing bovine serum albumin, glycosylated bovine serum albumin or advanced glycosylated endproduct of bovine serum albumin (This table also contains clones with no description in Examples).

In the table, EC\_G\_B/EC\_BSA and EC\_A\_B/EC\_BSA represent ratios of EC\_glycated\_BSA/EC\_BSA and EC\_AGE\_BSA/EC\_BSA, respectively.

			EC_glycated_	_BSA	EC_G_B	EC_A_B
10	Clone_name	EC_BSA	1	EC_AGE_BSA	/EC_BSA	/EC_BSA
	GAPDH(Cr1)	100.81	134. 21	115.16	1. 33	1.14
	βactin(Cr2)	1101.9	1092.57	997. 36	0.99	0. 91
15					_	
	ADRGL1000005	26.88	38. 27	36. 13	1	1
	ADRGL1000007	117. 89	127. <b>2</b> 5	133. 21	1.08	1. 13
	ADRGL1000009	29. 18	25. 65	26.05	1	1
20	ADRGL1000011	88. 9	117. 33	142. 9	1. 32	1. 61
	ADRGL1000027	33. 24	40. 53	43.02	1.01	1.08
	ADRGL1000058	153. 41	208. 84	180.05	1.36	1. 17
	ADRGL1000069	16.8	21. 77	29. 81	. 1	1
25	ADRGL1000077	25.74	24. 72	32. 86	1	1
<del></del>	ADRGL1000092	84. 52	84. 15	121. 76	1	1. 44
	ADRGL1000099	76.19	91.53	106. 01	1.2	1. 39
	ADRGL1000136	52. 34	44. 76	63.06	0.86	1. 2
30	ADRGL1000147	46.08	45. 18	52. 15	0.98	1. 13
30	ADRGL1000159	31.52	40. 24	42. 72	1.01	1.07
	ADRGL1000160	52. 34	60. 37	62. 29	1. 15	1. 19
	ADRGL1000171	21.46	16. 78	25. 59	1	1
	ADRGL1000181	37. 44	45. 71	43. 65	1. 14	1.09
35	B6G111000015	52. 42	71	65. 47	1.35	1. 25
	BGG111000016	127. 44	122. 93	147. 57	0.96	1. 16
	BGG111000017	25. 65	25, 74	31.33	1	1
	BGG111000022	32. 82	35. 19	25. 56	1	1
40	BGG111000031	44. 42	43. 8	40. 25	0. 99	0. 91
	BGG111000042	120. 38	146, 44	165, 42	1. 22	1. 37
	BGG111000046	74. 72	58, 85	84. 95	0. 79	1. 14
	BNGH41000020	4286.0	3584. 67	4330. 96	0.84	1. 01
45	BNGH41000025	216. 67	223. 74	257. 06	1.03	1. 19
	BNGH41000026	25. 76	28. 16	35. 52	1	1
	BNGH41000027	29. 23	23. 83	17.86	1	1
	BNGH41000035		238. 34	305. 66	0. 85	1.09
50	BNGH41000037	59. 14	54. 86	54. 58	0. 93	0. 92
	BNGH41000042	356. 1	324. 08	411.07	0. 91	1. 15
	BNGH41000048	1201.3	869. 03	739. 91	0. 72	0. 62
	BNGH41000056	33. 94	31.4	40.01	1	1
55	BNGH41000087	77. 58	81. 76	91.07	1.05	1. 17
33	BNGH41000091	21. 05	21. 23	26.82	1	1

	BNGH41000157	81. 11	57. 28	77. 46	0.71	0. 95
	BNGH41000169	21.1	17. 59	22. 53	1	1
	BNGH41000181	63. 54	56. 92	70.08	0. 9	1.1
5	BNGH41000198		26.38	34. 37	· 1	1
	BNGH41000219			142.46	1.4	1. 24
•	BNGH41000229		139.85	138.03	1.07	1.06
	BNGH41000237		91.81	114. 21	0. 97	1.21
10	BNGH41000238		62. 41	30.4	1.56	1
	BNGH41000243		78. 6	84. 17	0.87	0. 94
	BNGH41000270		30. 92	20.81	1	1
	BRAWH1000004		38. 29	36. 26	0.91	0. 91
15	BRAWH1000018	28. 94	43. 05	31.38	1.08	1
	BRAWH1000021	33.89	36. 68	36. 1	1	1
	BRAWH1000027	26. 95	26. 48	21.83	1	1
	BRAWH1000029	24. 31	26. 41		1	1
22	BRAWH1000040	49. 28	49. 78	51.7	1.01	1.05
20	BRAWH1000050	1118.2	957. 41	645. 63	0.86	0. 58
	BRAWH1000051	21.81	14, 16	15. 84	1	1
	BRAWH1000060	902. 17	1021.85	719.09	. 1.13	0.8
	BRAWH1000075		36. 18	34. 01	0. 98	
25	BRAWH1000081					
	BRAWH1000084			444. 54	1, 15	0. 98
	BRAWH1000095				1	1
	BRAWH1000096	37. 75		37.8	1	1
30	BRAWH1000097				1.03	0. 74
	BRAWH1000100				0. 75	0. 96
	BRAWH1000101				1.04	1
	BRAWH1000104				0.88	0. 88
35	BRAWH1000107				1	1
	BRAWH1000110			198. 95	1.28	0. 98
	BRAWH1000111			54. 17	1	1. 04
	BRAWH1000135			48.5	0. 89	1.09
40	BRAWH1000190				0. 99	0. 97
	HEMBA1000005			118. 59	0. 93	1. 11
	HEMBA1000006			31. 42	1	1
	HEMBA1000012			733. 9	1.12	0. 79
45	HEMBA1000020		1001.42	782. 76	0.91	0. 71
45	HEMBA1000030	39. 47	59. 22	55. 57	1.48	1. 39
	HEMBA1000034	41.06	33. 37	49.9	0.97	1. 22
	HEMBA1000042	87. 29	119. 44	105. 36	1.37	1. 21
	HEMBA1000045	38. 05	29. 97	32. 61	1	1
50	HEMBA1000046	65. 78	48. 47	56.5	0. 74	0.86
	HEMBA1000705	27. 41	22. 4	26. 81	1	1 01
	HEMBA1000713	69.05	62. 08	69. 75	0. 9	1.01
	HEMBA1000718	93. 98	62. 97	81.6	0. 67	0. 87
55	HEMBA1000719	26. 13	21.62	25. 73	1	1

	HEMBA1000722 29.02	25. 27	32.96	1	1
	HEMBA1000726 117.55	153. 37	143.95	1. 3	1. 22
	HEMBA1000727 67.77	72. 78	74.89	1.07	1.11
5	HEMBA1000732 61.56	51.31	53.93	0.83	0.88
	HEMBA1000736 38.55	46.78	24.16	1. 17	1
	HEMBA1000743 32.49	36. 94	42.85	1	1.07
	HEMBA1000745 57.36	47. 79	52. 26	0.83	0.91
10	HEMBA1000747 22.29	18. 39	22.12	1	1
-	HEMBA1000748 36.46	28. 26	36.84	1	1
	HEMBA1000749 74.95	85.11	86.02	1.14	1.15
	HEMBA1000752 39.73	44. 73	54.37	1.12	1.36
15	HEMBA1000753 40.75	48. 69	46.01	1.19	1.13
13	HEMBA1000757 35.64	44. 44	40.91	1, 11	1.02
	HEMBA1000760 69.3	67.6	93.08	0. 98	1.34
	HEMBA1000769 54.76	52. 63	53.06	0.96	0. 97
	HEMBA1000773 20.96	13. 95	22.38	1	1
20	HEMBA1000774 116.55	94. 14	94.73	0.81	0.81
	HEMBA1000780 31.69	25. 83	27.67	1	1
	HEMBA1000783 28.5	36. 63	26. 55	, 1	1
	HEMBA1000791 99.64	107. 65	83. 39	1. 08	0.84
25	HEMBA1000793 91.87	96. 91	85. 78	1. 05	0. 93
	HEMBA1000802 23.17	14. 63	15. 44	1	1
	HEMBA1000813 253.32	254. 92	342. 09	1. 01	1. 35
	HEMBA1000817 37.46	38. 22	41.77	1	1.04
30	HEMBA1000822 34.66	27. 08	34. 38	1	1
	HEMBA1000827 68.99	39. 78	47. 79	0. 58	0. 69
	HEMBA1000833 25.58	25. 52	30.57	1	1
	HEMBA1000835 2111.6	1497. 9	1485. 01	0.71	0.7
35	HEMBA1000843 56.27	54. 43	55. 21	0. 97	0. 98 1
	HEMBA1000851 31.69	22. 56	26.81	1	0.89
	HEMBA1000852 56.96	54. 44	50. 55	0, 96 0, 9	0.89
	HEMBA1000867 48.67	43. 93	29.75	0.96	0. 82
40	HEMBA1000869 41.88	38. 24 40. 06	35. 91 36. 77	0. 90	0.92
	HEMBA1000870 43.48 HEMBA1000872 104.68	84. 86	91.41	0. 92	0. 87
	HEMBA1000875 48.86	58. 31	55. 26	1. 19	1. 13
	HEMBA1000876 67.93	78. 45	71.48	1. 15	1.05
45	HEMBA1000907 70.99	54. 09		0. 76	1.05
,,,	HEMBA1000908 23.07	15. 94	19. 03	1	1
	HEMBA1000910 32.2	27. 08	30. 5	1	1
	HEMBA1000918 111.32	116. 33	81.06	1. 05	0.73
50	HEMBA1000919 22.07	24. 98	22. 53	1	1
50	HEMBA1000934 26.39	32. 83	28. 03	1	1
	HEMBA1000935 46.11	38. 81	46. 64	0. 87	1.01
	HEMBA1000940 85.36	71.35	78. 33	0. 84	0.92
	HEMBA1000942 70.42	56. 83	79. 23	0. 81	1.13
55					

	HEMBA1000943	12.93	13. 15	10.61	1	1
	HEMBA1000946	29.7	42. 14	32. 79	1. 05	1
	HEMBA1000960		190. 99	180. 48	1	0.95
5	HEMBA1000962	46. 12	59. 9	54. 5	1. 3	1.18
	HEMBA1000968	33. 79	36. 55	39. 28	1	1
	HEMBA1000971	48. 24	49.1	51.91	1. 02	1.08
	HEMBA1000972	38. 3	40. 09	39. 66	1	1
10	HEMBA1000374		56. 82	49. 75	1.09	0.96
	HEMBA1000975	46. 25	43. 7	33. 71	094	0.86
	HEMBA1000979		70. 71	68. 55	1. 11	1. 07
	HEMBA1000973		86. 95	94. 68	1. 34	1.46
	HEMBA1000983		48. 15	51.7	0. 94	1.01
15			27. 02	30. 11	1	1
	HEMBA1000985		49. 72	49. 93	0. 92	0. <del>9</del> 3
	HEMBA1000986			50.43	0. 32	0. 97
	HEMBA1000991		42. 47		1	0.37
20	· HEMBA1001007		24. 05	28. 85		
	HEMBA1001008		17. 62	20. 17	1	1
	HEMBA1001009		18. 86	22. 39	1	1
	HEMBA1001014		164. 28	150. 39	1. 22	1.11
0.5	HEMBA1001017		39. 24	67. 82	1	1.7
25	HEMBA1001019		35. 97	39. 91	1	1
	HEMBA1001020		62. 13	66. 28	1. 15	1. 23
	HEMBA1001021		29. 34	40. 37	1	1.01
	HEMBA1001022	41. 99	46. 11	37. 15	1.1	0. 95
30	HEMBA1001024	15. 72	18. 6	18. 46	. 1	1
	HEMBA1001026	17. 94	15. 49	20.39	1	1
	HEMBA1001043	25. 75	20. 78	29. 85	1	1
	HEMBA1001051	259.05	315. 04	343. 57	1. 22	1.33
35	HEMBA1001052	26. 16	30.4	30, 36	1	1
55	HEMBA1001059	214. 68	230. 87	246.86	1.08	1.15
	HEMBA1001060		78. 09	75. 1	1. 15	1, 11
	HEMBA1001064		26, 4	32. 27	1	1
	HEMBA1001071		32. 81	30.6	1	1
40	HEMBA1001077		36. 52	45. 31	1	1. 13
	HEMBA1001078		158. 37	287, 22	0.76	1.37
	HEMBA1001080		97. 65	106.07	1.09	1.18
	HEMBA1001084		60. 01	63, 93	0. 96	1.02
45	HEMBA1001085		46. 69	53. 3	0. 85	0.97
	HEMBA1001088		43. 37	46. 47	0. 9	0. 96
	HEMBA1001093		39.8	55. 42	0. 81	1. 12
			21. 59	19. 98	1	1
	HEMBA1001094		23. 52	26. 87	1	1
50	HEMBA1001099				1	1
	HEMBA1001104		37. 22	32. 3		1. 43
	HEMBA1001109		194. 24	223. 99	1. 24	
	HEMBA1001114		357. 72	505.7	0. 86	1. 21
55	HEMBA1001121	34. 94	28. 76	29. 96	1	1

	HEMBA1001122 -67.22	71. 48 43. 07	1.06	0. 64
	HEMBA1001123 68.59	45. 79 49. 75	0. 67	0. 73
	HEMBA1001133 26.76	19, 02 27, 29	1	1
5	HEMBA1001137 40.48	30. 55 37. 31	0.99	0. 99
	HEMBA1001140 47.02	45. 97 47. 6	0.98	1. 01
	HEMBA1001144 289.44	310. 11 365. 12	1. 07	1. 26
	HEMBA1001145 1034.5	1260. 22 1691. 61	1.22	1.64
10	HEMBA1001158 66.38	61.98 48.42	0. 93	0. 73
	HEMBA1001172 53.49	53. 98 48. 67	1.01	0. 91
	HEMBA1001174 29.2	20.8 21.66	1	1
	HEMBA1001175 90.3	84. 25 78. 88	0. 93	0. 87
15	HEMBA1001182 2256.3	1929. 4 1478. 23	0.86	0.66
15 .	HEMBA1001184 24.69	17. 33 21. 37	1	1
	HEMBA1001192 23.55	23. 79 26. 37	1	1
	HEMBA1001196 74.6	73. 23 78. 22	0. 98	1. 05
	HEMBA1001197 1122.0	1102.8 1110.66	0. 98	0. 99
20	HEMBA1001208 31.89	35. 83 - 28. 91	_ 1	1
	HEMBA1001213 2340. 3	1977, 61 1520, 51	0. 85	0. 65
	HEMBA1001214 32.13	33. 1 31. 22	. 1	1
	HEMBA1001221 16.36	19. 33 16. 96	1	1
25	HEMBA1001225 16.5	18.96 18.59	1	1
	HEMBA1001226 184.05	160, 45 175, 94	0.87	0.96
	HEMBA1001228 47.7	51. 37 42. 2	1.08	0.88
	HEMBA1001229 2013.5	2065, 33 1644, 74	1.03	0.82
30	HEMBA1001235 127.3	117. 6 147. 13	0. 92	1.16
	HEMBA1001238 93.08	83. 89 64. 54	0.9	0.69
	HEMBA1001242 1928. 4	1561, 31 1343, 98	0. 81	0.7
	HEMBA1001247 84.13	78. 26 71. 14	0. 93	0. 85
25	HEMBA1001253 1657.8	1315. 21 940. 34	0. 79	0. 57
35	HEMBA1001257 40.62	28. 56 45. 16	0. 98	1. 11
	HEMBA1001261 41.19	36, 59 33, 36	0. 97	0. 97
	HEMBA1001262 28.33	20. 98 26. 6	1	1
• •	HEMBA1001265 63.91	58. 2 61. 44	0. 91	0. 96
40	HEMBA1001266 83.23	75. 57 83. 48	0. 91	1
	HEMBA1001269 116.72	87. 43 124. 08	0. 75	1.06
	HEMBA1001272 29.45	23. 23 24. 56	1	1
	HEMBA1001279 362.45	336. 61 230. 89	0. 93	0.64
45	HEMBA1001281 380.63	330. 72 267. 19	0. 87	0.7
	HEMBA1001286 184.88	156. 57 159. 52	0. 85	0.86
	HEMBA1001289 25.32	27. 22 27. 13	1	. 1
	HEMBA1001291 135.14	121.49 137.23	0. 9	1.02
50	HEMBA1001294 66.23	38, 38 63, 36	0.6	0. 96
	HEMBA1001296 66.45	43. 63 35. 03	0.66	0. 6
	HEMBA1001297 38.2	62, 28 59. 53	1.56	1.49
	HEMBA1001299 145.18	178.69 152.2	1. 23	1.05
66	HEMBA1001302 91.92	114. 12 122. 61	1, 24	1.33
55				•

	HEMBA1001303	50.14	53. 38	45. 68	1.06	0.91
	HEMBA1001306		82. 23	86.06	1.07	1.12
	HEMBA1001308	•	173. 73	151.85	1.07	0. 94
5	HEMBA1001310	35. 02	29. 23	32. 29	1	1
	HEMBA1001312		120. 24	124.38	1.04	1.08
	HEMBA1001319		35. 57	28. 57	1	1
	HEMBA1001322		31.63	29. 19	1	1
10	HEMBA1001323		29. 58	36.66	1	1
	HEMBA1001326		62.6	81.26	0.84	1.09
	HEMBA1001327		52. 98	44. 45	1. 32	1, 11
	HEMBA1001330		106. 44	132.86	0.84	1.05
15	HEMBA1001348		28.5	36. 52	1	1
13	HEMBA1001350		46.03	43.08	1, 11	1. 04
	HEMBA1001351		99,5	89. 31	0.89	0.8
	HEMBA1001352		54. 89		1.28	1. 29
	HEMBA1001353		440. 78	378.96	1. 15	0. 99
20	HEMBA1001358		148	158. 17	1.03	1.1
	HEMBA1001361		35. 45	46. 23	1	1. 16
	HEMBA1001364		25. 6	38. 53	. 1	1
	HEMBA1001375		50. 18	55. 42	1. 24	1. 36
25	HEMBA1001377	123.11	130. 48	138. 05	1.06	1. 12
	HEMBA1001383	24. 75	21.89	25. 34	1	1
	HEMBA1001387	7 133. 99	110. 07	119. 25	0.82	0. 89
	HEMBA1001388	16.86	15. 81	19. 96	1	1
30	HEMBA1001390	162. 95	166. 12	229. 96	1.02	1. 41
	HEMBA1001391		25. 43	34. 84	1	1
	HEMBA1001398	3 71.07	92. 62	94. 61	1.3	1. 33
	HEMBA100140	5 18.28	22. 16	27. 71	1	1
35	HEMBA100140	6 47.72	56. 62	52. 03	1. 19	1. 09
	HEMBA100140	7 26.06	28. 51	42.35	1	1.06
	HEMBA100141		23. 93	33. 22	1	1
	HEMBA100141	3 26.86	28. 89	35. 64	. 1	1
40	HEMBA100141		88. 49	97. 78	1. 67	1. 85
40	HEMBA100141		56. 61	61. 78	1.31	1. 43
	HEMBA100141		70. 23	83. 44	1.01	1. 21
	HEMBA100143		46. 16	57. 09	1.07	1. 33
	HEMBA100143		93. 43	95. 11	1. 18	1.2
45	HEMBA100143					1. 21
	HEMBA100144	_	18. 75	24. 89	1	1 22
	HEMBA100144		99.2	116.86	1. 13	1. 33
	HEMBA100145		60. 97	62. 25	0. 82	0. 84
50	HEMBA100145		101. 22	82. 71	1, 17	0. 95
	HEMBA100145		33.8	38. 01	1	1 00
	HEMBA100145		36. 67	43. 17	1	1. 08
	HEMBA100146		38. 47	33. 2	1	1
55	HEMBA100146	29.08	29. 65	31. 99	1	

	HEMBA1001463	59. 9	67.79	69: 08	1.13	1. 15
	HEMBA1001469	70.99	49.11	46. 76	0.69	0.66
	HEMBA1001473 1		148.99	167.:31	0.88	0.99
5	HEMBA1001477	24. 92	25. 29	23. 86	1	1
•	HEMBA1001478		24. 36	30. 89	1	1
	HEMBA1001480		56. 83	53. 79	1, 05	1
			52. 6	67. 18	0. 9	1.14
10	HEMBA1001490		34. 25	39. 52	1	1
10	HEMBA1001495		1323.57	1231.86	1. 25	1.17
	HEMBA1001497		73. 45	74. 91	1.09	1.11
	HEMBA1001510		193. 22	183.1	1.18	1. 12
	HEMBA1001515		50. 63	34. 96	1.23	0.97
15	HEMBA1001517		108.99	110. 39	0.97	0.98
	HEMBA1001522		30.46	22.06	1	1
	HEMBA1001526		49. 14	54.08	1.03	1. 13
	HEMBA1001533		71.35	81.79	0.94	1.07
20	HEMBA1001547		84.8	90. 33	1.03	11
	HEMBA1001552		124. 65	127. 65	0.83	0.85
	HEMBA1001553		480.55	526. 67	1.04	1.13
	HEMBA1001557		94.47	76.06	1.02	0.82
25	HEMBA1001563		59.53	45. 05	0. 94	0.71
	HEMBA1001566		88. 26	79. 39	1.07	0.96
	HEMBA1001569	128. 2	118. 39	122. 68	0. 92	0. 96
	HEMBA1001570	155. 44	162.32	161. 01	1.04	1.04
30	HEMBA1001579	56. 65	97. 72	71. 55	1. 72	1. 26
	HEMBA1001581		159. 23	140. 29	1. 39	1. 22
	HEMBA1001582	41.6	30. 97	50. 61	0.96	1. 22
	HEMBA1001585		54. 87	46. 66	1. 15	0. 97
35	HEMBA1001589		47. 65	37. 92	1.14	0.96
	HEMBA1001595		45. 3	48. 48	0. 72	0. 77 1
	HEMBA1001604	28. 63	29. 02	33. 97	1	
	HEMBA1001608		155. 85	151. 37	0. 84 1. 23	0. 82 1. 27
40	HEMBA1001615		926. 26	957. 18	1.23	1, 27
40	HEMBA1001620		147. 63	174. 25 24. 38	1.03	1, 21
	HEMBA1001621	26. 51	20. 21		0. 82	0. 73
	HEMBA1001635		45. 05 20. 65		1	1
	HEMBA1001636		74. 45		0. 86	1. 24
45	HEMBA1001640		2251.04	2043. 18	0. 93	0. 84
	HEMBA1001647		110. 51	108. 89	1. 15	1. 13
	HEMBA1001651 HEMBA1001655		77. 88		1. 08	1. 01
	HEMBA1001658		33. 83		0. 78	0. 83
50	HEMBA1001661		27. 02		1	1
	HEMBA1001665		33. 84		1	1
	HEMBA1001670		69. 39		0. 82	1. 16
	HEMBA1001672		36. 31		1	1
55	HEMDATOCIO/2		22. 21			

	HEMBA1001673	82. 64	87. 42	58.92	1,06	0. 71
	HEMBA1001675		36. 24	36.07	1	1
	HEMBA1001676		567. 16	633. 55	1. 13	1. 27
	HEMBA1001678		282, 33	287. 87	0. 97	0. 99
	HEMBA1001680	59	71.89	67. 27	1. 22	1. 14
	HEMBA1001681		53. 31	69.16	1. 21	1.58
	HEMBA1001684		101.34	128. 16	0.83	1. 05
	HEMBA1001695		28. 18	28. 49	· 1	1
	HEMBA1001702		36. 11	34. 21	1	1
	HEMBA1001709		23. 23	24. 21	1	1
	HEMBA1001711		49. 57	49. 1	1. 2	1. 19
15	HEMBA1001712	22. 12	21. 69	26. 49	1	1
13	HEMBA1001714		150. 33	176.72	1.09	1. 28
	HEMBA1001717		28. 98	29. 74	1	1
	HEMBA1001718	108. 75	108. 53	111.93	1	1.03
	HEMBA1001723	86. 07	96. 41	63, 93	1.12	0. 74
20	HEMBA1001731	<b>26.03</b> .	27. 17	29. 94	1	1
	HEMBA1001734	60.16	66. 91	55, 53	1. 11	0. 92
	HEMBA1001736	48. 23	61. 31	61.56	. 1.27	1. 28
	HEMBA1001741	43. 58	39. 35	44. 77	0. 92	1.03
25	HEMBA1001744	23. 18	17. 43	22. 95	1,	1
	HEMBA1001745	18. 96	20. 05	34. 02	1	1
	HEMBA1001746		60. 61	65. 09	1, 01	1.08
	HEMBA1001761		48. 07	52. 97	1. 16	1. 28
30	HEMBA1001762		82. 14	101. 34	0. 94	1. 16
	HEMBA1001781	39. 61	36. 99	39. 24	1	1
	HEMBA1001784		54. 9	68. 72	1. 17	1.47
	HEMBA1001791	46. 46	66. 14	73. 08	1. 42	1.57
35	HEMBA1001794		133.9	156.75	1. 26	1.48
	HEMBA1001800		1898. 36	1564. 1	1. 18	0. 97
	HEMBA1001803	25. 79	35. 11	28. 19	1	1
	HEMBA1001804		169. 79	195. 66	1.19	1. 37
40	HEMBA1001808		33. 25	41.14	. 1	1. 03 1. 03
	HEMBA1001809		79. 05	92. 5	0. 88	1. 03
	HEMBA1001811		100. 48	102. 21 56. 79	1. 09 1. 27	1. 28
	HEMBA1001815		56. 35 19. 93	29. 04	1.27	1. 28
45	HEMBA1001816		65. 42	75. 42	0. 89	1. 03
	HEMBA1001819 HEMBA1001820	73. 56 19. 68	22. 61	21. 88	1	1. 00
	HEMBA1001822		40. 45	37. 76	1. 01	1
	HEMBA1001824		187. 12	202. 26	1. 1	1. 19
	HEMBA1001835		29. 39	36. 01	0. 99	0. 99
50	HEMBA1001844		89. 37	81. 98	1. 12	1.03
	HEMBA1001847		29. 06	36. 37	1. 12	1.00
	HEMBA1001849		116. 04	94. 33	1. 49	1. 21
	HEMBA1001850		147. 28	101.43	1. 03	0.71
55			==			

	HEMBA1001861	26. 36	23.37	20.95	1	1.
	HEMBA1001862		506.65	561.78	0.97	1.08
	HEMBA1001864		36.34	50. 44	1	1. 26 ·
5	HEMBA1001866		103.03	113.87	1. 16	1.29
	HEMBA1001869		67. 41	64. 56	1.03	0.99
	HEMBA1001871		492. 28	509.64	0. 85	0.88
	HEMBA1001876		129. 25	159. 26	0. 82	1.01
10	HEMBA1001878		47.78	44. 75	1.05	0. 98
70	HEMBA1001879		34.3	29.06	1	1
	HEMBA1001884		38. 01	37.87	0. 97	0. 97
	HEMBA1001886		48.62	51.77	0.98	1.04
	HEMBA1001888		124. 27	117.94	1.12	1.06
15	HEMBA1001890		43. 36	39. 58	1.01	0. 93
	HEMBA1001896		28. 53	28.11	1	1
	HEMBA1001899		458. 41	380. 57	1.02	0.85
	HEMBA1001904		2362	2247.7	0. 95	0.9
20	HEMBA1001910		21.39	18.7	1	1
	HEMBA1001911	102.05	84. 21	89.81	0. 83	0.88
	HEMBA1001912		615.96	496.77	0. 82	0.66
	HEMBA1001913		69.81	74. 33	1. 23	1. 31
25	HEMBA1001915	26.54	25. 2	30. 07	1	1
	HEMBA1001918	66.46	78. 62	61. 54	1. 18	0. 93
	HEMBA1001921	37.68	40. 38	33. 39	1.01	1
	HEMBA1001931	17. 26	20. 51	14.88	1	1
30	HEMBA1001939		29. 42	29. 03	1	1
	HEMBA1001940		55. 22	56. 02	0. 82	0. 83
	HEMBA1001942		19. 95	24. 59	1	1
	HEMBA1001944		917. 16	769, 79	1.08	0. 91
35	HEMBA100194		29. 26	33.14	1	1
	HEMBA1001950		37.1	38. 34	1	1 0. 86
	HEMBA100195		149. 14	144. 1	0. 89 1	0.86
	HEMBA100195		28. 04	28. 27	0.88	1.06
40	HEMBA100196		104	125. 36 20. 74	0.00	1.00
40	HEMBA100196		21. 92 36. 48	40. 09	0. 97	0. 97
	HEMBA100196			58. 17	1, 1	1.03
	HEMBA100196		61. 72 31. 5	32. 33	1. 1	1. 00
	HEMBA100197			158. 33		1
45	HEMBA100198		119. 27	118. 39	1.03	1.02
	HEMBA100199 HEMBA100200		124. 44	174. 33	0.86	1. 2
			82. 27	75, 78	0.86	0.79
	HEMBA100200		93. 59	67. 04	1.01	0.72
50	HEMBA100200 HEMBA100201		31.56	29. 3	1	1
	HEMBA100201		33. 11	28. 77	1	1
	HEMBA100202		788. 12	855.8	0. 69	0.75
	HEMBA100203		35. 38	36. 56	1	1
55	TILIIDA TOCCO		22.30			

	HEMBA1002035 32.58	39. 5	34. 54	1	1
	HEMBA1002037 35.32	25. 71	34. 06	1	1
	HEMBA1002038 80.78	95. 24	87. 43	1.18	1.08
5	HEMBA1002039 29.27	41.94	45. 05	1.05	1.13
•	HEMBA1002042 69.56	82. 26	75. 06	1. 18	1.08
	HEMBA1002043 94.56	132.79	125. 69	1.4	1.33
	HEMBA1002048 27.06	23. 59	27. 16	1	1
10	HEMBA1002049 98. 28	115.71	107. 64	1. 18	1.1
10	HEMBA1002053 109.41	96.4	115. 83	0.88	1.06
	HEMBA1002055 94.18	84. 55	102. 03	0.9	1.08
	HEMBA1002056 36.57	41. 32	37. 72	1.03	1
	HEMBA1002061 30.02	37. 92	40. 57	1	1.01
15	HEMBA1002080 217.45	341.32	408. 08	1. 57	1.88
	HEMBA1002084 25.19	26. 67	28. 24	1	1
	HEMBA1002085 39.74	35. 38	59. 04	1	1.48
	HEMBA1002092 25.03	22. 15	28	1	1
20	HEMBA1002098 29.75	28. 25	35. 17	1	1
	HEMBA1002100 290.2	250. 1	284. 9	0.86	0. 98
	HEMBA1002101 141.62	117. 14	110	0. 83	0. 78
	HEMBA1002102 63.92	79. 33	78. 63	1. 24	1. 23
25	HEMBA1002105 104.76	105. 75	87. 22	1. 01	0.83
	HEMBA1002107 651.41	734. 58	1023. 32	1. 13	1. 57
	HEMBA1002113 209.87	180. 63	238. 46	0.86	1.14
	HEMBA1002119 783.32	765. 1	693. 78	0. 98	0. 89
30	HEMBA1002125 81.76	58. 98	98. 89	0.72	1. 21
	HEMBA1002131 61.47	44. 91	63	0.73	1. 02
	HEMBA1002133 41.34	51.09	47. 54	1. 24	1. 15
	HEMBA1002139 22.23	20. 8	27. 64	1	1
35	HEMBA1002141 30.33	26. 59	38. 58	1	1
	HEMBA1002144 62.71	82. 97	95. 04	1.32	1.52
	HEMBA1002147 55. 36	91. 25	90. 72	1.65	1. 64
	HEMBA1002150 14.67	23. 86	28. 41	1	1
	HEMBA1002151 99.61	91. 89	115. 21	0. 92	1. 16 1. 15
40	HEMBA1002153 58.68	76. 71		1. 31	
	HEMBA1002156 14.27	16.44		1 45	1 1. 67
	HEMBA1002160 54.01	78. 26	90. 28	1. 45 1. 13	1. 07
	HEMBA1002161 170.98	192.46	221. 4 72. 69		
45	HEMBA1002162 52.85	75. 43		1. 12	1. 34
	HEMBA1002163 110.56	123. 56	148. 01 766. 76	1. 14	1. 54
	HEMBA1002164 765.65	871.05		0. 91	1. 22
	HEMBA1002166 549. 48	498. 57	670. 97 36. 06	0. 31	1. 22
50	HEMBA1002167 38.8	26.87	49. 35	1.08	0. 99
	HEMBA1002173 49.98	53. 97 26. 26	30. 98	1.00	0. 33
	HEMBA1002177 28.1	26. 26 38. 21	56. 1	0. 91	1. 27
	HEMBA1002178 44.04	38, 21 300, 94	273. 96	1. 24	1. 13
55	HEMBA1002179 242.19	300. 84	4/0. 70	1. 47	0

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	HEMBA1002185		180. 38	169. 67		1
	HEMBA1002188		55. 72	64.06		1. 28
	HEMBA1002189		109.04	134. 1		1. 43
5	HEMBA1002191	72.79	83. 6	75. 92		1.04
	HEMBA1002192	42	46.84	42. 55	1. 16 1. 15 1. 12 1. 03 0. 99 1. 01 1. 0. 91 1. 0. 94 1. 18 1. 1. 0. 96 1. 05 0. 95 0. 89 0. 84 1. 1 1. 18 0. 97 7. 1 1. 1. 18 0. 97 7. 1 1. 01 1. 08 0. 72 1. 17 1. 03 0. 89 6. 1 7. 1 1. 03	1.01
	HEMBA1002195	54. 39	56. 23	65. 6		1. 21
	HEMBA1002196	44. 2	43. 54	63. 36	0. 99	1.43
10	HEMBA1002199	37. 95	40. 26	40.09	1.01	1
•	HEMBA1002204	23. 94	31.87	26. 05	1	1
	HEMBA1002208	441.74	403.49	389. 44	0. 91	0.88
	HEMBA1002212		25. 82	32. 44	1	1
45	HEMBA1002215		39. 95	40. 56	0. 94	0. 95
15	HEMBA1002217		1497. 14	1240. 47	1.18	0.98
	HEMBA1002220		21. 45	25. 27	1	1
	HEMBA1002226		236. 81	242	1. 04	1.06
	HEMBA1002227		357. 19	334. 86	1.1	1.03
20	HEMBA1002229		103. 52	123.73	0. 96	1. 15
	HEMBA1002237		82. 87	75. 28	1. 05	0.96
	HEMBA1002239		2194. 9	1725. 53	0.95	0. 75
	HEMBA1002241		754. 66	596. 7	0.89	0.7
25	HEMBA1002253		33.51	38. 9	0. 84	0.84
	HEMBA1002257		24. 24	20.65	1	1
	HEMBA1002259		17. 9	26. 65	1	1
	HEMBA1002262		202. 87	176. 45	1. 19	1.04
30	HEMBA1002265		31.84	33. 14	1	1
••	HEMBA1002267		70.88	58. 1	1. 18	0. 97
	HEMBA1002270	80. 52	78.06	58. 36	0. 97	0.72
	HEMBA1002286		21.23	21. 47	1	1
	HEMBA1002290	39. 01	40. 25	44. 87	1. 01	1.12
35	HEMBA1002302		184. 71	149. 25	1.08	0.87
	HEMBA1002304		57. 92	62. 29	0. 72	0.78
	HEMBA1002307		402. 44	441.62	1. 17	1.29
	HEMBA1002316	43. 33	44. 78	43. 11	1.03	0.99
40	HEMBA1002319	79.54	70.87	88. 85	0.89	1.12
	HEMBA1002320	31. 21	22. 68	27. 16	1	1
	HEMBA1002321	33.5	29. 95	24. 57	1	1
	HEMBA1002328	31.99	44.02	42.84	1.1	1.07
45	HEMBA1002333	27. 85	26. 55	26. 56	1	1
	HEMBA1002337		96. 97	92. 69	1.05	1
	HEMBA1002339		3357. 21	2424.94	1.09	0.79
	HEMBA1002341		25. 97	35.7	1	1
50	HEMBA1002348		50. 98	51. 42	1.12	1.12
	HEMBA1002349	23.97	27. 43	26. 69	1	1
	HEMBA1002353		43. 17	43. 08	0.9	0.9
	HEMBA1002356		151.54	132. 62	1.1	0.97
	HEMBA1002357		3477.73	2585.96	1.29	0.96
55						

	HEMBA1002360 1	19. 97	102.96 ··	109.34	0. 86	0.91
		39. 83	33. 85	40.98	1	1.02
	HEMBA1002365		34. 97	39.04	0.93	0.93
5		24. 04	18.89	25. 62	1	1
	HEMBA1002374		49. 95	80.12	0. 71	1.15
	HEMBA1002376 2		2235. 67	2311. 2	0.82	0.85
	HEMBA1002377 1		168.08	175. 4	1.28	1.33
10	HEMBA1002380		120. 28	112.72	1.14	1.07
70		90. 89	94. 67	100.47	1.04	1.11
	HEMBA1002384	25. 99	33. 81	24.8	1	1
			61.03	39.69	0. 93	0.61
15	HEMBA1002396		92.86	105. 59	1.01	1. 15
15	HEMBA1002402		27. 12	36.05	1	1
	HEMBA1002417		23. 18	39. 92	1	1
	HEMBA1002419	23. 6	33. Ó7	24. 23	1	1
	HEMBA1002420		108.83	115. 26	1.04	1. 1
20	HEMBA1002421	25. 47	33. 19	32. 71	1	- 1
	HEMBA1002423	30. 15		35. 32	1	1
	HEMBA1002424		50. 34	58. <b>57</b>		1.21
	HEMBA1002426	95. 14	82. 18	130. 83	0.86	1.38
25	HEMBA1002430		22. 86	48. 75	1	1. 22
	HEMBA1002439		39. 45	35. 77	0.87	0.87
	HEMBA1002441		151. 58	198.51	0.8	1.04
	HEMBA1002454	18. 11	16.69	21. 15	1	1 57
30	HEMBA1002458	89. 76	106. 49	140. 61	1.19	1. 57 1. 01
		48. 27	40. 03	48. 58	0.83	1. 49
		62. 66	51.68	9306	0. 82 1	1.49
	HEMBA1002465	24. 06	23. 59	34. 77 878	0. 97	0. 77
35	HEMBA1002469		1105. 08 45. 86		0.9	1. 03
	HEMBA1002475	50.7	89. 51		1. 23	1. 23
	HEMBA1002477	72. 48 66. 92	53. 09		0. 79	
		88. 24	115. 58		1.31	
40	HEMBA1002481 HEMBA1002486		70. 33	89. 87	1.07	1. 37
	HEMBA1002490		55. 01		0.89	0. 97
	HEMBA1002495		28. 23	37, 46	1	1
	HEMBA1002498		39. 98	50. 12	1	1. 25
45	HEMBA1002501				1.16	1. 15
	HEMBA1002503		65. 1	76.49	1.08	1. 27
	HEMBA1002504		92. 18	96. 39	1.29	1.35
	HEMBA1002508	•	97. 62	141. 37	1.01	1. 47
50	HEMBA1002513		20. 42	33. 54	1	1
	HEMBA1002515		28. 92	24.09	1	1
	HEMBA1002524		29. 99	28. 95	1	1
	HEMBA1002538		33. 86	28.82	1	1
	HEMBA1002542		85. 68	127, 33	0. 85	1. 27
55						

	HEMBA1002544 36.36	43. 78	37. 89	1.09	1
	HEMBA1002546 233.93	235. 61	280. 15	1.01	1.2
	HEMBA1002547 104.87	101.89	134. 44	0.97	1. 28
5	HEMBA1002550 77.5	74.04	63. 14	0.96	0.81
	HEMBA1002551 32.81	34.15	38.69	1	1
	HEMBA1002552 68.67	77.66	73.63	1. 13	1. 07
	HEMBA1002555 18.78	18. 95	18. 73	1	1
10	HEMBA1002558 89.84	96.83	105. 37	1. 08	1. 17
70	HEMBA1002561 66.15	70, 22	75. 71	1.06	1.14
	HEMBA1002562 18.32	22.04	22. 77	1	1
	HEMBA1002568 37.87	37.42	38. 86	1	1
	HEMBA1002569 42.87	36. 63	53. 23	0. 93	1. 24
15	HEMBA1002570 77.85	65. 44	95. 45	0.84	1. 23
	HEMBA1002574 24.93	18.6	20, 86	1	1
	HEMBA1002583 53.05	51. 72	50. 08	0. 97	0.94
	HEMBA1002587 156. 82	157. 42	134. 58	1	0.86
20	HEMBA1002590 128. 54	122.14	121. 34	0. 95	0. 94
		75. 31	82. 11	0. 98	1.06
	HEMBA1002592 77.18 HEMBA1002595 31.08	27. 57	36. 21	1	1
	HEMBA1002609 700.66	946. 91	778.73	1. 35	1.11
25	HEMBA1002617 108.51	91.58	109.76	0.84	1. 01
23	HEMBA1002619 53.7	51.89	47. 88	0. 97	0. 89
	HEMBA1002621 25.77	20. 09	21.51	1	1
	HEMBA1002624 122.93	107. 08	101.57	0. 87	0. 83
	HEMBA1002628 46. 21	31.91	41. 12	0. 87	0. 89
30	HEMBA1002629 52.6	81.02	53. 39	1. 54	1. 02
	HEMBA1002632 113.01	97.63	105. 71	0. 86	0. 94
	HEMBA1002645 91.47	117. 47	109. 31	1. 28	1.2
	HEMBA1002651 44.04	49. 34	47. 34	1. 12	1.07
35	HEMBA1002652 57. 48	51.49	60. 52	0. 9	1.05
	HEMBA1002659 71.71	69. 66	66. 98	0. 97	0. 93
	HEMBA1002661 45.57	46. 97	41.7	1. 03	0. 92
	HEMBA1002666 31.14	34. 85	20. 48	1	1
40	HEMBA1002667 22.1	24. 84	22. 87	1	1
	HEMBA1002673 159.38	123.39	164. 21	0.77	1.03
	HEMBA1002678 72.5	72. 15	68. 16	1	0.94
	HEMBA1002679 28.83	30. 33	32. 02	1	1
	HEMBA1002688 83.9	60. 71		0. 72	1
45	HEMBA1002696 40.11	28. 05	35. 95	1	1
	HEMBA1002703 88.57	94.66	125. 63	1.07	1, 42
	HEMBA1002705 88.37	53. 4	47. 13	1.34	1. 18
•		141. 19	141. 71	1. 04	1.04
50	HEMBA1002712 135.74 HEMBA1002715 1896.2	141. 19	1150. 31	0. 78	0. 61
		33.83	36. 74	0. 73	1
	HEMBA1002716 34.19 HEMBA1002718 160.42	97. 63	151. 44	0. 61	0. 94
	HEMBA1002718 100. 42	105. 61	106. 33	1.11	1. 12
55	HEIRDA I 002/20 93. 30	100.01	100. 55	•••	

	HEMBA1002730	100.98	92.09	106.69	0. 91	1.06
	HEMBA1002734		55. 32	63.59	0.87	1
	HEMBA1002742		33, 17	32.33	1	· 1
5	HEMBA1002746		31. 17	37.84	1	1
	HEMBA1002748		48. 28	41.23	1.08	0. 92
	HEMBA1002750		77. 01	7 <b>7</b>	1.16	1.16
	HEMBA1002755		94. 45	103. 73	0. 97	1.07
	HEMBA1002759		40. 64	57. 32	1. 02	1.43
10	HEMBA1002763		1337. 96	1262.67	0. 79	0. 74
	HEMBA1002767	_	43. 14	69.82	0.77	1. 24
	HEMBA1002768		42.89	38. 86	1.07	1
	HEMBA1002769		32. 49	43. 28	1	1.08
15	HEMBA1002770		93	93. 35	1. 18	1. 19
	HEMBA1002777		32. 84	36. 1	1	1
	HEMBA1002779		88. 2	99. 28	1.1	1.24
	HEMBA1002780		91.7	97. 2	1.03	1.09
20	HEMBA1002790		77. 32	81.02	0.99	1.04
	HEMBA1002794		30. 2	40.09	1	1
	HEMBA1002798		43. 12	47. 67	0.96	1.06
	HEMBA1002801		30.11	32.1	1	1
25	HEMBA1002810		55.01	71.45	1. 13	1.46
	HEMBA100281		34. 48	41. 11	1	1, 03
	HEMBA1002811		1785. 51	1207.17	1.1	0.74
	HEMBA1002820		106. 18	120. 59	1. 03	1.17
30	HEMBA100282		37. 96	64. 29	0. 86	1.38
30	HEMBA100283		125. 28	100. 54	1. 17	0.94
	HEMBA100285	0 24.01	25. 95	26. 09	1	1
	HEMBA100286	2 240.02	315. 35	291. 51	1.31	1.21
	HEMBA100286	3 44. 25	38. 73	38. 24	0.9	0. 9
35	HEMBA100286	7 33.51	29. 59	29. 49	1	1
	HEMBA100287	6 55.95	60. 55	78. 99	1.08	1, 41
	HEMBA100288	6 23.5	21. 75	42. 15	1	1. 05
	HEMBA100289	6 52.27	44. 86	64. 45	0.86	1. 23
40	HEMBA100291		51.85	67.01	1. 18	1, 53
	HEMBA100292	1 19.64	28. 3	24.8	1	1
	HEMBA100292	4 39.54	32. 67	39. 68	1	1
	HEMBA100293		245. 83		1. 13	1, 34
45	HEMBA100293	85 64.16	64. 74	108. 58	1.01	1.69
	HEMBA100293		35.04	<b>62. 25</b>	1	1.56
	HEMBA100293		34. 91	34. 91	1	1
	HEMBA100294		57. 33	79. 04	1.43	1. 98
50	HEMBA100295				1	1
30	HEMBA100295	54 35.46			1	1
	HEMBA100290		_		1.05	1.34
	HEMBA10029				1.41	1. 42
	HEMBA10029	70 149.32	307. 3	233. 48	2.06	1.56

	HEMBA1002971 28.	82 44. 02	38. 42	1.1	1
	HEMBA1002973 110.		152. 71	1. 31	1. 38
	HEMBA1002978 16.		27. 46	1	1
5	HEMBA1002981 24.		29.89	1	1
	HEMBA1002985 31.		46. 14	1	1. 15
	HEMBA1002986 57.		65. 26	0.96	1. 13
	HEMBA1002988 31.	•	50. 95	1	1. 27
	HEMBA1002992 278		353. 67	1. 26	1. 27
10	HEMBA1002995 63.		70. 57	1. 02	1.12
	HEMBA1002997 27.		41.04	. 1	1.03
		45 18.81	22. 48	1	1
	· · · · · · · · · · · · · · · · · · ·	72 14. 99	19.67	1	1
15		.56 39.38	47. 56	0. 94	1. 12
•		. 59 33. 5	43.5	1	1.09
	HEMBA1003021 140	1	158.62	1. 14	1.13
		. 32 59. 67	62.18	1.1	1.14
20	HEMBA1003029 350		345.82	0. 87	0. 99
		. 91 89. 75	89. 48	0.9	0.9
		. 33 54. 76	58. 64	1.01	1.08
	HEMBA1003033 115		133.35	1. 28	1.16
25		7. 5 178. 84	195. 02	1.3	1.42
25	• • • • • • • • • • • • • • • • • • • •	. 36 18. 2	25. 53	1	1
		2. 07 37. 83	48.71	0.77	0.94
	HEMBA1003041 142		165.95	1.11	1.17
	HEMBA1003046 191		201.43	1.03	1.05
30		3. 52 70. 12	63.88	1.1	1.01
		3. 24 27. 22	36. 82	1	1
		1. 27 75. 36	69. 57	0.83	0.76
		52. 1 51. 43	· 56.27	0. 99	1.08
35	HEMBA1003071 39	9. 64 57. 64	58. 19	1.44	1. 45
	HEMBA1003072 3:	3. 87 32. 7	35. 83	1	1
	HEMBA1003076 26	2. 68 221. 2	263. 78	0. 84	1
	HEMBA1003077 2	1. 44 19. 93	32.74	1	1
40	HEMBA1003078 4	1. 19 40. 76	37. 58	0. 99	0. 97
	***************************************	7. 77 101. 97	97.73	1. 04	1
	HEMBA1003083 1	13.8 121.46	124. 26	1.07	1.09
	• • • • • • • • • • • • • • • • • • • •	2. 19 50. 66	57. 16	0. 97	1.1
45	HEMBA1003090 10		107. 55	0. 79	1.01
	***	2. 18 25. 87	27. 93	1	1
		8. 21 31. 54	41. 6	1	1.04
	***************************************	79. 4 81. 64	100.06	1.03	1. 26
	••	8. 08 38. 19	54. 34	0. 83	1.13
50		24. 92 25. 81	24. 36	1	1 10
		51. 66 56. 64		1, 1	1.19
	HEMBA1003117	33. 2 28. 07		1	0.93
	HEMBA1003120 6	6. 51 67. 4	55. 48	1. 01	0. 83

	HEMBA1003129 171.02	190. 86	212. 79	1.12	1. 24
	HEMBA1003133 64.48	64. 84	68. 91	1.01	1.07
	HEMBA1003136 28.75	29. 95	35. 68	1	1
5	HEMBA1003142 85.8	71. 23	84.93	0.83	0.99
	HEMBA1003148 21.58	18. 05	30. 73	1	1
	HEMBA1003151 44.39	33. 7	45. 39	0.9	1.02
	HEMBA1003152 73.95	101.12	37. 82	1.37	0.54
10	HEMBA1003157 23.61	44.7	43, 82	1.12	1.1
,,	HEMBA1003166 396.48	495. 89	465.55	1. 25	1. 17
	HEMBA1003171 24.29		38. 03	1	1
	HEMBA1003175 91.24		99.96	1.2	1.1
	HEMBA1003179 511.42	505.09	456.99	0.99	0.89
15	HEMBA1003186 130.84		160.21	0.88	1. 22
•	HEMBA1003196 60.73		68.65	1. 14	1.13
	HEMBA1003197 23.09	32. 24	20.62	1	1
	HEMBA1003199 30.15	33. 48	35. 67	1	1
20	HEMBA1003202 160.12	213. 03	163.11	1. 33	1.02
	HEMBA1003204 82.45	94, 23	93, 26	1. 14	1.13
	HEMBA1003210 172.76	185. 07	241.07	1.07	1.4
	HEMBA1003212 116.79	142. 68	125.64	1. 22	1.08
25	HEMBA1003218 24.25		25. 18	1.04	1
	HEMBA1003220 773.18		891. 78	0. 97	1. 15
	HEMBA1003222 28.08		35. 63	1	1
	HEMBA1003225 12.05		15. 46	1	1
30	HEMBA1003229 19.34		31. 73	1	1
	HEMBA1003230 29.03		35. 38	1	1
	HEMBA1003235 83.94		89. 7	1. 32	1.07
	HEMBA1003236 116. 94		123	0.8	1.05 1
35	HEMBA1003250 19. 2		26. 24	1 0. 95	0. 98
	HEMBA1003252 223.		219. 85 119. 67	1. 34	1.3
	HEMBA1003257 92.00 HEMBA1003268 28.20		39, 23	1. 04	1. 3
			57. 42	1, 15	1.21
	HEMBA1003273 47.5 HEMBA1003276 46.4		55. 78	0. 96	1, 2
40	HEMBA1003277 10.0		17. 36	1	1
	HEMBA1003278 28.3		38. 31	1	1
	HEMBA1003280 44.2		53. 2	0. 91	1. 2
	HEMBA1003281 21.5		27. 7	1	1
45	HEMBA1003284 24.9		26. 49	1	1
	HEMBA1003286 717.8		607.04	1.3	0.85
	HEMBA1003291 38.9		55. 08	1. 13	1.38
	HEMBA1003294 54.2		71. 14	1, 15	1.31
50	HEMBA1003296 1300.		1115. 39	1.09	0.86
	HEMBA1003304 22.8		27. 12	1	1
	HEMBA1003306 49.1		68. 74	1. 14	1.4
	HEMBA1003309 25.0	9 28.4	32. 7	1	1

	HEMBA1003314 37.17	44. 82	59. 15	1. 12	1.48
	HEMBA1003315 124.59	124. 67	175. 23	1	1.41
	HEMBA1003312 56. 36	76.18	78. 78	1. 3 <b>5</b>	1.4
5	HEMBA1003326 30.66	36. 54	33. 24	1	1
3	HEMBA1003327 36.18	42. 14	47. 09	1. 05	1. 18
		49. 8	59. 69	1. 25	1. 49
		95. 56	101. 26	1. 15	1. 22
	HEMBA1003330 83.32	290. 07	281. 18	1. 27	1. 23
10	HEMBA1003348 228.39	34. 25	41.3	1. 27	1. 03
	HEMBA1003369 35.08	127. 15	160.35	1. 12	1. 42
	HEMBA1003370 113.05	41. 81	33. 29	1. 05	1
	HEMBA1003373 30.18	175. 71	168.37	1, 16	1. 11
15	HEMBA1003376 151.06	31. 1	35. 25	1. 10	1
	HEMBA1003380 23.37	29. 28	28. 6	1	1
	HEMBA1003384 23.97	18. 54	15. 1	1	1
	HEMBA1003387 14.31		37. 16	1	1
20	HEMBA1003392 26.1	18. 97 46. 06	38. 3	1. 15	1
	HEMBA1003395 35.41	23. 91	34. 6	1. 13	1
	HEMBA1003399 22.92	47. 83	55.33	0. 97	1. 12
	HEMBA1003400 49. 2	22. 88	27. 46	1	1
	HEMBA1003402 19.93 HEMBA1003403 916.83	1093. 91	744. 48	1. 19	0. 81
25		32.87	29.66	1	1
	HEMBA1003408 31.78 HEMBA1003412 104.21	93. 23	112. 18	0. 89	1. 08
	HEMBA1003417 38.97	45. 45	46. 16	1. 14	1. 15
	HEMBA1003417 33.57	157. 73	180.06	1. 17	1. 34
30	HEMBA1003410 134. 33		205. 39	0. 87	1, 12
	HEMBA1003425 23.51	25. 24	25. 02	1	1
	HEMBA1003433 19.55		22. 1	1	1
	HEMBA1003440 18.36		18. 41	1	1
35	HEMBA1003442 118. 4		106. 66	0. 82	0.9
	HEMBA1003447 1244. 4		857.98	1.08	0.69
	HEMBA1003453 95.87		110.27	0. 97	1. 15
	HEMBA1003461 26.09		34. 23	1	1
40	HEMBA1003463 32.85		36.19	1.02	1
	HEMBA1003465 32.65		39. 45	1	1
	HEMBA1003480 150.99		160.32	1.21	1.06
	HEMBA1003485 60.29		46. 93	0. 98	0.78
	HEMBA1003487 41.67		45. 62	0. 96	1.09
45	HEMBA1003492 50.94		58. 23	1	1.14
	HEMBA1003494 81.07		63.81	0. 79	0. 79
	HEMBA1003497 20.15		23. 08	1	1
	HEMBA1003503 21.56		27. 4	1	1
50	HEMBA1003511 24.64		33. 32	1	1
	HEMBA1003528 30.56		32. 98	1	1
	HEMBA1003530 21.5		27. 5	1	1
	HEMBA1003531 78.03		79. 44	0.96	1.02

	HEMBA1003532 97.65	102. 85	110. 25	1.05	1.13
	HEMBA1003538 103.67	110.46	109.77	1.07	1.06
	HEMBA1003545 40.23	24. 96		0.99	0.99
5	HEMBA1003546 26.04	26. 92		1	1
-	HEMBA1003548 108.9	123. 89		1.14	1
	HEMBA1003553 46. 21	39. 71	96 21.54 0.99 0.99 92 35.27 1 89 109.3 1.14 71 61.93 0.87 1. 77 39.61 1 61 59.09 0.81 1. 62 174.51 1.15 1. 27 284.97 0.97 1. 68 22.4 1 94 62.36 1.27 1. 3.3 86.06 1.05 1. 44 38.45 1 07 30.17 1 0.2 21.85 1 .77 115.09 0.84 0. 42 25.76 1.04 .48 80.25 1.18 178 113.04 1.16 153 109.44 1.08 179 34.55 1 .03 56.92 1.04 084 38.91 1.05 .48 83.33 0.93 1 .22 103.55 1.18 1 .42 26.9 1 .44 24.57 1 .503 63.5 1.05 1 .609 109.82 0.97 1	1.34	
	HEMBA1003555 26. 42	35. 77		1	1
	HEMBA1003556 54.88	44, 61		0.81	1.08
10	HEMBA1003560 166.71	191. 62	174. 51	1.15	1.05
	HEMBA1003565 201.56	195. 27	284. 97	0.97	1.41
	HEMBA1003568 24.9	29.68	22. 4	1	1
	HEMBA1003569 39.46	50.94	62.36	1. 27	1.56
15	HEMBA1003571 75.65	79.3	86.06	1.05	1.14
	HEMBA1003579 25.07	31. 44	38. 45	1	1
	HEMBA1003580 26.85	27. 07	30.17	1	1
	HEMBA1003581 23.83	19. 2	21.85	1	1
20	HEMBA1003591 119.29	100.77	115.09	0.84	0.96
	HEMBA1003595 24.19	41.42	25. 76	1.04	1
	HEMBA1003597 65.92	77. 48	80. 25	1.18	1.22
	HEMBA1003598 20.14	18.68	21. 51		1
25	HEMBA1003600 100.74	116. 78	113.04		1. 12
	HEMBA1003602 89.14	96. 53	109.44		1. 23
	HEMBA1003604 27.67	30. 92			1
	HEMBA1003610 28.27	27. 79			1
30	HEMBA1003615 58.86	61.03			0. 97
	HEMBA1003617 26.21	41. 84			1
	HEMBA1003620 65.82	61.48			1. 27
	HEMBA1003621 85.87	101. 22			1.21
	HEMBA1003622 28.46	22. 42			1
35	HEMBA1003630 21.72	20. 46			
	HEMBA1003637 43.95	46. 03			1. 44 1. 07
	HEMBA1003640 102.34	99.09		0.97	1.07
	HEMBA1003645 22.81	24. 65	19. 83 31. 99	1	1
40	HEMBA1003646 23.59	29. 71	20. 76	i	1
	HEMBA1003647 15.69	19. 27 161. 88	151.69	1. 68	1. 57
	HEMBA1003656 96.49	27. 04	45. 76	1. 00	1. 14
	HEMBA1003662 30.59 HEMBA1003666 10.75	13. 44	15.76	1	1
45		59. 07	108. 02	0. 81	1.48
	HEMBA1003667 73.07 HEMBA1003670 19.08	13, 39	19.91	1	1
	•	295. 13	431.7	1.02	1.49
	HEMBA1003674 288.8 HEMBA1003677 55.17	79. 59	79. 49	1. 44	1.44
50		65. 84	77. 2	1.05	1. 23
	HEMBA1003679 62.99 HEMBA1003680 93.24	87. 1	126. 5	0. 93	1.36
	HEMBA1003684 24.94	28. 67	31. 31	1	1
	HEMBA1003690 92.14	94. 42		1.02	1. 21

	HEMBA1003692 58.18	64. 91	87. 68	1.12	1.51.
	HEMBA1003702 29.67	40. 2	41.34	1.01	1.03
	HEMBA1003711 32.3	33. 55	41.53	1	1.04
5	HEMBA1003714 11.66	12.85	14, 26	1	1
	HEMBA1003715 127. 28	128. 58	176. 49	1.01	1.39
	HEMBA1003717 36.23	47. 11	56.82	1.18	1.42
	HEMBA1003720 72.7	104.3	95.71	1. 43	1.32
10	HEMBA1003725 30. 21	29.52	45. 16	1	1.13
	HEMBA1003728 19.89	22.33	35.98	1	1
	HEMBA1003729 34.19	28. 4	38. 87	1	1
	HEMBA1003732 24.57	23.52	26.8	1	1
	HEMBA1003733 32.05	38. 41	35. 69	1	1
15	HEMBA1003742 91.11	74.8	121.1	0. 82	1.33
	HEMBA1003743 20.87	27.7	36. 19	1	1
	HEMBA1003758 155.66	212.5	192.43	1.37	1.24
	HEMBA1003760 19.4	24. 15	21.44	1	1
20	HEMBA1003764 50.94	59.98	66.6	1. 18	1.31
	HEMBA1003769 243.2	236. 31	294. 05	0. 97	1.21
	HEMBA1003773 23.79	30. 65	30. 94	. 1	1
	HEMBA1003783 46.84	51.35	61.57	1.1	1.31
25	HEMBA1003784 23.33	24. 87	32.65	1	1
	HEMBA1003794 98.42	151.82	160. 49	1.54	1.63
	HEMBA1003799 24	20. 43	26.88	. 1	1
	HEMBA1003803 92.09	92. 45	94. 87	1	1.03
30	HEMBA1003804 11.79	13. 4	13.71	1	1
	HEMBA1003805 62.9	47. 84	57.71	0. 76	0. 92
	HEMBA1003807 35	33. 04	37. 16	1	1
	HEMBA1003810 49.79	42.58	48. 65	0.86	0. 98
35	HEMBA1003827 214.33	220. 34	247. 88	1. 03	1. 16
	HEMBA1003836 105.3	131.09	152.63	1. 24	1.45
	HEMBA1003838 201.86	193. 21	218. 25	0.96	1.08
	HEMBA1003843 95.01	80. 4	81, 33	0.85	0. 86 0. 9
•	HEMBA1003846 584.93	521. 93	524. 6	0.89	0.9
40	HEMBA1003856 37.85	19. 35	24. 84	1 02	1.1
	HEMBA1003857 84.45	85. 97	93	1. 02 1. 06	1. 15
	HEMBA1003864 45.81	48. 61	52. 54	1.00	1. 13
•	HEMBA1003866 22.19	14. 38	26. 76 171. 8	1. 21	1. 35
45	HEMBA1003868 127. 43	154. 52 75. 5	61. 32	1. 24	1. 01
	HEMBA1003879 60. 94		59. 16	1. 16	0. 93
	HEMBA1003880 63.83	74. 3 265. 18	244. 1	0. 9	0. 83
	HEMBA1003884 295.83	77. 54	85, 49	1	1.1
50	HEMBA1003885 77.57	77. 54 44. 46	51. 02	1. 03	1. 18
	HEMBA1003887 43.2	30. 68	46. 62	1.03	1. 17
	HEMBA1003890 34.02	90. 01	86. 46	1	0. 96
	HEMBA1003893 89.77	108.77	130. 61	1. 53	1.84
	HEMBA1003896 70.89	100.77	100.01		

	HEMBA1003902 - 72.1	82. 48	88. 57	1. 14	1. 23
	HEMBA1003904 26.68	30. 31	35. 19	1	1
	HEMBA1003908 -20.58	24. 77	17.24	1	1
5	HEMBA1003926 817.82	708.16	810.39	0. 87	0. 99
	HEMBA1003937 74.16	92. 69	106. 26	1. 25	1. 43
	HEMBA1003939 189.88	194. 21	169.09	1. 02	0.89
	HEMBA1003940 27. 59	32. 67	41.23	1	1.03
10	HEMBA1003941 39.11	38. 63	32.44	1	1
10	HEMBA1003942 29.31	36. 49	35. 17	1	1
	HEMBA1003945 101.81	111.03	153.5	1.09	1.51
	HEMBA1003949 26.89	23. 59	31.98	1	1
	HEMBA1003950 108. 43	101.22	111.88	0. 93	1.03
15	HEMBA1003953 33.64	26. 45	19.18	1	1
	HEMBA1003958 172. 27	318. 81	393.92	1.85	2. 29
	HEMBA1003959 37.17	45. 76	40. 45	1.14	1.01
	HEMBA1003960 41.6	37.03	46.77	0.96	1.12
20	HEMBA1003966 50.41	56. 91	- 67. 77	1.13	1.34
	HEMBA1003967 35.26	32.76	36.11	1	1
	HEMBA1003968 17.83	15. 7	30.29	, 1	1
	HEMBA1003974 5265.0	4839.83	4912.1	0.92	0.93
25	HEMBA1003976 20.51	33. 59	27. 16	1	1
	HEMBA1003977 23.04	26.84	23. 25	1	1
	HEMBA1003978 37.27	32.98	40. 15	1	1
	HEMBA1003981 108	97.04	117. 19	0. 9	1.09
30	HEMBA1003982 1665.9	1749.33	1773. 93	1.05	1.06
50	HEMBA1003985 26.77	25. 32	36. 45	1	1
	HEMBA1003987 33.45	37. 68	41. 63	1. 19 0. 9 1. 93 1. 05 1. 45 1	1.04
	HEMBA1003989 63.17	62.72	76. 11	0. 99	1.2
	HEMBA1004000 58.48	80.61	79. 89	1.38	1. 37
35	HEMBA1004006 88.99	136.49	109.66	1.53	1.23
	HEMBA1004007 98.97	105.35		1.06	1.05
	HEMBA1004010 777.44	800.96	1113.06	1.03	1. 43
	HEMBA1004011 23.51	25. 26	29. 93	1	1
40	HEMBA1004012 55.61	56.63	63. 69	1.02	1. 15
	HEMBA1004015 67.35	53.48		0.79	1. 03
	HEMBA1004024 98.84	120.03		1.21	1. 28
•	HEMBA1004029 23.09	21.98	27. 27	1	1
45	HEMBA1004038 23.15	29. 19	29. 88	1	1
	HEMBA1004042 20.59	28. 59	29. 34	1	1
	HEMBA1004045 30	25. 74	41. 36	1	1.03
	HEMBA1004048 87.48	104. 2	129. 71	1. 19	1. 48
50	HEMBA1004049 46.55	56. 7	79. 44	1.22	1. 71
<b>V</b> V	HEMBA1004051 24.85	23, 39	30. 94	1	1
	HEMBA1004053 83.24	94. 23		1.13	1.48
. ,	HEMBA1004055 21.51	22. 83		1	1
	HEMBA1004056 118.22	132. 94	128. 78	1.12	1.09
55					